

54679_g_at	inhibitor of kappa light polypeptide gene enhancer in B-cells, kinase beta	IKBKB	AA521493	2.2	1.5	3.9
63839_at	CGI-24 protein		AA923141	2.1	1	3.9
50252_at			AA149410	3.9	2.9	3.9
72644_at			AA938448	3.5	2.3	3.9
48106_at	hypothetical protein FLJ20489		H14241	4.1	3.6	3.9
38487_at	KIAA0246 protein		D87433	3.8	3.6	3.9
55564_at	spleen tyrosine kinase	SYK	AA521157	3.9	3.8	3.9
55664_at	annexin A11	ANXA11	AI923271	3.2	1.9	3.9
54948_at	HSPC128 protein		AI684747	4.1	3.5	3.9
55598_at	KIAA0692 protein		AI188279	3.3	3.6	3.8
39520_at	ubiquitously transcribed tetratricopeptide repeat gene, Y	UTY	AI924382	0.9	1.4	3.8
34477_at	chromosome		AF000994	5.6	2.4	3.8
37475_at	DKFZP434J046 protein	SEPW1	AC004144	0.5	0.9	3.8
39709_at	selenoprotein W, 1	ITGA5	U67171	2.6	1.4	3.8
39753_at	integrin, alpha 5 (fibronectin receptor, alpha polypeptide)	CSNK2A1	X06256	1.9	3.1	3.8
594_s_at	casein kinase 2, alpha 1 polypeptide		M55265	0.9	1.7	3.8
40619_at	ubiquitin carrier protein	CALM1	M91670	2.8	7.2	3.8
955_at	calmodulin 1 (phosphorylase kinase, delta)	NAP1L1	M27319	1.1	0.7	3.8
40634_at	nucleosome assembly protein 1-like 1		M86667	0.4	0.5	3.8
33309_at			AA521060	0.1	0.3	3.8
33869_at	trinucleotide repeat containing 15	TNRC15	AL080218	5.2	3.2	3.8
35785_at	nuclear receptor subfamily 1, group D, member 1	NR1D1	W28281	16.4	3.3	3.8
32520_at	FabG (beta-ketoacyl-[acyl-carrier-protein] reductase, E coli) like	FABGL	M24900	1.5	0.5	3.8
41440_at			D82061	1.7	1.6	3.8
799_at	cyclin-dependent kinase 5, regulatory subunit 1 (p35)	CDK5R1	X80343	1	0.3	3.8
36281_at	KIAA0478 gene product		M96739	0.4	0.2	3.8
35990_at	SEC22, vesicle trafficking protein (S. cerevisiae)-like 1	SEC22L1	AB007947	5.9	1.1	3.8
41598_at	protein kinase, Y-linked	PRKY	AA890010	4	2.2	3.8
40030_at	solute carrier family 5 (sodium/glucose cotransporter), member 2	SLC5A2	Y15801	0	0.8	3.8
41305_at	ATPase, Na+/K+ transporting, beta 3 polypeptide	ATP1B3	M95549	1.2	1.6	3.8
32563_at			U51478	6.4	3.6	3.8

Figure 13IIIII

37363_at	KIAA0429 gene product		AB007889	7.9	7.6	3.8
37922_at	transcobalamin II; macrocytic anemia	TCN2	L02648	0.9	3.2	3.8
38764_at	helicase-moi		AF007142	0.2	0	3.8
33805_at	KIAA0480 gene product		AB007949	9	2.5	3.8
39091_at	vitamin A responsive; cytoskeleton related		AF070523	0.5	1.6	3.8
34416_at	Cas-Br-M (murine) ecotropic retroviral transforming sequence	CBL	X57110	2.3	0	3.8
58005_at	hypothetical protein		AI480253	3.7	6.7	3.8
48497_at	membrane-associated nucleic acid binding protein		AI380524	3.5	2	3.8
66326_at			AI936888	9.5	5	3.8
76071_at			AA031499	1.1	1.2	3.8
46511_at			R38554	0.4	1.7	3.8
76724_r_at			H38425	0.2	1	3.8
55019_at	hypothetical protein FLJ21610		R61377	0.1	2.7	3.8
59934_at			AI684648	1.2	0.7	3.8
82177_at			H94893	4.5	0	3.8
78106_at			AI634844	0.1	0	3.8
90840_at			W88900	0.8	0.3	3.8
82273_at			AW014723	4	2.1	3.8
79388_at			AI128130	0.9	0.8	3.8
47890_at			AI921465	1.6	0.7	3.8
54926_at			R53602	2	1	3.8
80313_at			AI215686	3	1.9	3.8
63189_at			AI342495	1.4	0.6	3.8
84492_at			AI288213	0	0.9	3.8
54807_at			N72621	0.4	3.4	3.8
85581_at			AI332404	2.6	1.8	3.8
48566_at	hypothetical protein FLJ20446		N98615	7.7	6.4	3.8
85100_at			AI567916	5.7	3	3.8
43092_at			AI925903	3.6	2	3.8
56292_at			R98956	1.4	3.4	3.8
56903_r_at			T97939	8.6	0.4	3.8
69763_r_at			AI865825	4.9	0.9	3.8
70691_at			AI631466	2.3	0.8	3.8
42851_g_at			AI884619	1.4	1.5	3.8
48463_at			AA362254	2.5	4.4	3.8

Figure 13JJJJ

49626_at	Alg5, <i>S. cerevisiae</i> , homolog of		AA425251	18.6	4.7	3.8
51035_f_at	hypothetical protein FLJ22672		AA310453	3.4	3.9	3.8
44832_at			R26252	0	0	3.8
53821_at			AI075913	2.5	2.6	3.8
44115_at	hypothetical protein from EUROMAGE 588495		AA486504	27.1	3.1	3.8
58356_at		EPB49	AA142978	3.9	2.3	3.8
75162_r_at	erythrocyte membrane protein band 4.9 (dema1in)		AI673236	1	0.5	3.8
49299_s_at	mitochondrial carrier homolog 1		AA449775	0.9	1.1	3.8
61641_at			AA479481	4	1.5	3.8
91341_at			AI291659	5.9	4.5	3.8
54532_at	melanocortin 1 receptor (alpha melanocyte stimulating hormone receptor)	MC1R	AI123000	4.1	2.4	3.8
58434_at	zinc finger protein 146	ZNF146	AI268312	6.1	2.4	3.8
58734_at			N48694	0	0	3.8
53604_at			AI928164	1.6	2.2	3.8
46874_at			AA699559	1.1	1.2	3.8
64422_at			AA634254	3.6	2.3	3.8
46911_at			AI381767	1	0.3	3.8
64457_at	leucine rich repeat (in FLII) interacting protein 2	LRRFIP2	AA203144	8.6	4.4	3.8
55809_at			AA028139	10.5	3.3	3.8
47106_at	hypothetical protein MGC3169		AA196034	0.2	0.4	3.8
45488_s_at			AI525989	0.9	0.5	3.8
887_at	growth differentiation factor 1	GDF1	M62302	0.8	0.3	3.8
55481_i_at			R54660	0.3	0.2	3.8
61500_f_at			N62149	1.4	2.1	3.8
86349_at			T70214	5.8	2.7	3.8
88335_r_at	WD repeat domain 7	WDR7	AA126277	0.9	0.6	3.8
41430_at	mitogen-activated protein kinase kinase kinase 5	MAP3K5	AB011113	4	1.2	3.8
1327_s_at	nuclear factor of activated T-cells, cytoplasmic, calcineurin-dependent 3	NFATC3	U67156	1.4	1.6	3.8
40822_at			L41067	5.3	4.4	3.8
41349_at	presenilin 2 (Alzheimer disease 4)	PSEN2	L43964	1.9	1.1	3.8
36647_at	hypothetical protein FLJ10326		AA526812	3.5	1.7	3.8
34839_at	KIAA1104 protein		AB029027	1.4	0.9	3.8
33920_at	diaphanous (Drosophila, homolog) 1	DIAPH1	AF051782	4.3	3.2	3.8

Figure 13KKKKK

198_g_at	non-metastatic cells 3, protein expressed in	NME3	U29656	3.4	2.7	3.8
38780_at	aldo-keto reductase family 1, member A1 (aldehyde reductase)	AKR1A1	J04794	3.5	0.8	3.8
48087_at	ubiquitin-conjugating enzyme E2E 2 (homologous to yeast UBC4/5)	UBE2E2	AI432317	2.6	3.1	3.8
61287_f_at	similar to rat nuclear ubiquitous casein kinase 2		AI380204	3.1	2.2	3.8
76484_at			AA789296	3.8	3.4	3.8
63374_at	mucoilin 1	MCOLN1	AI816064	3.2	1.5	3.8
65701_at	hypothetical protein FLJ20254		AI554392	3.2	1	3.8
67162_r_at			AI366683	3.8	3.3	3.8
48524_at			AI659343	0.2	1.3	3.8
84668_at	LENG1 protein		AI701895	2.6	2.1	3.8
44743_at			AF062725	3.1	0.8	3.8
65561_at			AL120093	2.8	0	3.8
45249_at			AA775778	3.6	2.2	3.8
48806_at	N-terminal kinase-like	NTKL	AI741642	1.8	1.4	3.8
43382_s_at	C-terminal binding protein 1	CTBP1	AI991853	3.4	2.5	3.8
64314_s_at			AI653557	3.6	3.4	3.8
58213_at			AI926429	0.5	1.8	3.8
44343_s_at	hypothetical protein, estradiol-induced		AI074877	3.9	3.1	3.8
61116_g_at			AA442301	2.7	2.3	3.8
46569_at			AA552976	3.8	2.8	3.8
46233_at	hypothetical protein FLJ20420		AI341488	2.6	3.3	3.8
48952_at	hypothetical protein FLJ21634		AI985274	3.8	3.1	3.8
68967_at	serine/threonine kinase 22B (spermiogenesis associated)	STK22B	AA573033	3.8	2.3	3.8
39534_at	lysosomal-associated protein transmembrane 4 alpha		L77564	1.3	0.5	3.7
39019_at	endothelial differentiation, lysophosphatidic acid G-protein-coupled receptor, 4	EDG4	D14696	1	1.5	3.7
37769_at			AF011466	1.7	0.8	3.7
34902_at	KIAA0492 protein	EFNA2	AB007961	0.1	0	3.7
33090_at	ephrin-A2		AJ007292	2.1	0.5	3.7
40980_at	KIAA1564 protein		W26477	1.1	1.8	3.7
41156_g_at	catenin (cadherin-associated protein), alpha 1 (102kD)	CTNNA1	U03100	9.1	5.3	3.7
1959_at	antizyme inhibitor		D88674	2.5	3	3.7
41282_s_at	peroxisome biogenesis factor 10	PEX10	AA194159	0.5	0.4	3.7

Figure 13LLLLL

35138_at	A kinase (PRKA) anchor protein 8	AKAP8	Y11997	0	0.1	3.7
39582_at			AL050166	1.6	1.2	3.7
40595_at	Treacher Collins-Franceschetti syndrome 1	TCOF1	AI345337	1.5	0.3	3.7
37517_at	KIAA1039 protein	HSPA6	AB028962	1.9	1.1	3.7
117_at	heat shock 70kD protein 6 (HSP70B)		X51757	2.6	0.6	3.7
34808_at	KIAA0999 protein		AB023216	6.5	1	3.7
41808_at	cyclin D binding Myb-like transcription factor 1	DMTF	AF052102	2.8	1.9	3.7
36580_at	integrin-linked kinase	ILK	U40282	1	4.7	3.7
37374_at	hypothetical protein FLJ13910	ANXA4	AL050139	0	0.7	3.7
36580_at	annexin A4	BGN	M82809	0	3	3.7
38126_at	biglycan	GABRB1	J04599	0	0	3.7
35028_at	gamma-aminobutyric acid (GABA) A receptor, beta 1		X14767	0	0	3.7
43059_at	KIAA0955 protein		AI040349	2.7	1.1	3.7
57734_s_at			AA677558	0	0	3.7
61471_at			N66393	6	4.9	3.7
58192_at	KIAA1411 protein		AI308839	1.3	0.8	3.7
46493_at			N99350	3.7	0.8	3.7
42355_r_at			AA699719	6.2	2.9	3.7
48116_at			AI096389	1.8	2.5	3.7
58769_at			AA604596	2.9	3.2	3.7
56959_at			AI376649	2.7	2.8	3.7
57008_at	cortic al thymocyte receptor (X. laevis CTX) like		AI799005	0.7	0.7	3.7
75557_at			AI446519	1.2	0.7	3.7
66303_i_at			R43631	1.7	0.8	3.7
90309_at			AI076012	2.6	2.6	3.7
76805_at			AI332830	2.9	1.6	3.7
66839_at			AI138572	2.5	1.5	3.7
77643_f_at			AA502152	2.4	0.9	3.7
77855_at			AI435014	1.8	2.1	3.7
65199_at			AI078337	6.5	10	3.7
48929_at			AA521311	3.1	2.5	3.7
42584_at			AI571796	0.3	2.3	3.7
82731_f_at			AI792894	1	0.7	3.7
52763_at			AI277429	20.2	4.3	3.7
91350_at			AA989283			

Figure 13MMMMMM

84119_at			AI939467	1.1	0.5	3.7
60041_at	transmembrane 4 superfamily member (tetraspan NET-2)		AI654147	2	1.6	3.7
91865_at			AI417837	3.3	2.3	3.7
64962_r_at	poly(A)-binding protein, cytoplasmic 1	PABPC1	AI356950	0.4	1.9	3.7
43217_at			AA642418	2.5	1.3	3.7
59765_at			H17909	2.9	2.6	3.7
45240_at	stromal cell protein		N93439	0.5	0.6	3.7
44409_r_at	ribosomal protein L13	RPL13	N51496	2.1	1.5	3.7
67141_at			AI267607	6.6	1.1	3.7
86377_at			N63995	2.5	2.9	3.7
55018_at	hypothetical protein FLJ12517		W46342	0.5	1.2	3.7
86606_at			AA884702	0.9	3.8	3.7
68364_at			AI074578	0.7	1.6	3.7
75254_at	matrilin 2	MATN2	AI972357	0.8	0	3.7
62794_at	hypothetical protein FLJ10989		AI861854	1.7	2.1	3.7
71309_at			AI919139	0.3	0.9	3.7
71504_at			AA811476	1.8	1.2	3.7
59632_at			AI991777	0.1	1	3.7
72080_at			N99035	0	0.3	3.7
72778_at			AI655731	0.2	4.2	3.7
64262_at			R39949	0.9	0.6	3.7
45483_at			AI253027	1.9	1.5	3.7
50933_at	Alport syndrome, mental retardation, midface hypoplasia and elliptocytosis chromosomal region, gene 1	AMMECR1	AL037664	0	0.1	3.7
66148_i_at			AW000952	1	6	3.7
44089_at			W26293	2.1	5.3	3.7
50587_at	trichorhinophalangeal syndrome 1	TRPS1	AA024604	1.5	1.9	3.7
42052_at	sialic acid binding Ig-like lectin 7	SIGLEC7	AA400618	1	2.1	3.7
57757_r_at			AA425516	0	0	3.7
80081_at	ribosomal protein L4	RPL4	AI721208	0.5	0.5	3.7
44631_at			AI653257	0.2	3.1	3.7
83244_r_at			AI308504	1.4	2	3.7
47406_at			R50967	1.8	1.6	3.7
72742_at			AA250992	3.6	3.3	3.7
55971_at	KIAA1323 protein		AI263115	0	0	3.7

Figure 13NNNNN

51708_at	KIAA0495			AI971075	1.5	1	3.7
55166_at				AA593942	1	1.3	3.7
64714_at				AI828075	0.3	1.9	3.7
64466_at	RNA binding motif protein 7		RBM7	AI990688	4.3	2.5	3.7
54477_at				AI193993	0	1.1	3.7
45348_at	hypothetical protein			N32178	2.5	1.5	3.7
46286_s_at	hypothetical protein FLJ20080			AA534905	1.2	0.8	3.7
62572_at				AI140917	0	3.4	3.7
44609_at	KIAA1254 protein			AA001419	0.7	1.3	3.7
68127_at				AA308556	0	0	3.7
54461_at				AI250784	0.7	2	3.7
54478_at				AI638525	2.3	1.6	3.7
86420_at				H30740	2.6	1	3.7
85179_at				AL041520	0	1.1	3.7
40279_at	KIAA0121 gene product		SLC25A6	D50911	5.1	3.4	3.7
40435_at	solute carrier family 25 (mitochondrial carrier; adenine nucleotide translocator), member 6			J03592	2.9	1.8	3.7
32853_at	translocase of outer mitochondrial membrane 70 (yeast) homolog		TOMM70A	AB018262	2.5	1.4	3.7
36013_at	chromosome 4 open reading frame 1		C4ORF1	AF006621	2.6	1.4	3.7
36185_at	alanyl-tRNA synthetase		AARS	D32050	4.3	1.3	3.7
40695_at	IMP (inosine monophosphate) dehydrogenase 1		IMPDH1	J05272	2.9	2.7	3.7
39155_at	proteasome (prosome, macropain) 26S subunit, non-ATPase, 3		PSMD3	D67025	2.5	1.6	3.7
48084_at				AA582193	3.7	3.2	3.7
52896_at	translocase of inner mitochondrial membrane 10 (yeast) homolog		TIMM10	AI498204	2.2	3	3.7
56529_at	APMCF1 protein			AA156214	3.3	3.6	3.7
90257_at	hypothetical protein DKFZp547D065			AI089323	3.6	2.6	3.7
51223_at	phosphoinositide-3-kinase, catalytic, beta polypeptide		PIK3CB	AI916796	4.1	1.8	3.7
60062_s_at	hypothetical protein FLJ13556 similar to N-myc downstream regulated 3			AI027140	3.5	2.9	3.7
43857_s_at				AA570070	3.5	2.3	3.7
65645_at	IGF-II mRNA-binding protein 2			AI675886	3.1	1.8	3.7
49181_at				AI192291	1.9	1	3.7

Figure 13000000

61712_at	hypothetical protein PR00823		AI990512	2.8	2.1	3.7
55678_at	hypothetical protein MGC10986		W72885	3.6	3.2	3.7
45805_g_at	hypothetical protein FLJ20643		AA948226	3.1	2.9	3.7
64443_at	EH-domain containing 4	EHD4	AI971203	1.2	0.8	3.7
47390_at			AA928060	3.7	3.7	3.7
64502_s_at	hypothetical protein FLJ20859		AW006969	3.3	1.5	3.7
46220_at	peroxisomal short-chain alcohol dehydrogenase		AI131178	3.3	3	3.7
54112_at			AA443434	4.4	2.3	3.7
54935_at	CD74 antigen (invariant polypeptide of major histocompatibility complex, class II antigen-associated)	CD74	N22918	4	3.2	3.7
49228_s_at			AW026208	2.2	2.2	3.7
44536_at			AA523300	3.2	1.8	3.7
37459_at	collagen, type VIII, alpha 1	COL8A1	X57527	0.2	0	3.6
39404_s_at	regulator of nonsense transcripts 1	RENT1	D86988	1.3	1.4	3.6
41871_at	lung type-I cell membrane-associated glycoprotein		AI660929	20.8	0	3.6
39694_at	hypothetical protein MGC5508		W27517	0	0.5	3.6
31867_at			AF052174	1.8	5.2	3.6
32142_at	serine/threonine kinase 3 (Ste20, yeast homolog)	STK3	U26424	5	4.1	3.6
40864_at	ras-related C3 botulinum toxin substrate 1 (rho family, small GTP binding protein Rac1)	RAC1	D25274	5.3	2.5	3.6
41147_at			AF038186	4.1	15.4	3.6
32379_f_at	pyruvate kinase, muscle	PKM2	W28740	1.2	1.6	3.6
34182_at			U18932	1.6	1.6	3.6
35738_at	high-mobility group (nonhistone chromosomal) protein 17-like 3	HMG17L3	AI347088	6.9	3.9	3.6
32720_at	Apg12 (autophagy 12, S. cerevisiae)-like	APG12L	AA151716	2.3	2.3	3.6
32905_s_at	tryptase, alpha	TPS1	M30038	3.8	2.3	3.6
830_at	Tax1 (human T-cell leukemia virus type I) binding protein 1	TAX1BP2	U25801	5.2	0.8	3.6
34846_at	calcium/calmodulin-dependent protein kinase (CaM kinase) II	CAMK2B	AF112472	3.6	2	3.6
	beta					
36452_at	synaptopodin		AB028952	4.5	1.6	3.6
36737_at	crystallin, beta A4	CRYBA4	U59057	3.9	0	3.6
35325_at	GTPase Rab14		AF052113	0.5	0.8	3.6
35826_at	suppressor of Ty (S.cerevisiae) 5 homolog	SUPT5H	AF040253	4.5	3.1	3.6
36918_at	guanylate cyclase 1, soluble, alpha 3	GUCY1A3	Y15723	18.2	3.1	3.6

Figure 13PPPPP

36987_at	lamin B2	LMNB2	M94362	1.7	1.1	3.6
38452_at			U90904	0	2	3.6
38453_at	intercellular adhesion molecule 2	ICAM2	X15606	2.5	0.3	3.6
1083_s_at	mucin 1, transmembrane	MUC1	M35093	0.1	0.2	3.6
46750_s_at	hypothetical protein PR02219		AA479496	2.1	3	3.6
56804_at			AA916789	12.3	14.3	3.6
50533_at			AI084504	0.5	1.1	3.6
49642_at	sorting nexin 2	SNX2	N59854	3.7	0.7	3.6
47421_at			AI701034	1.5	1.7	3.6
90174_r_at			AI218205	0.4	1.2	3.6
57379_at			AA865085	1	0.6	3.6
42524_at			R37337	30.1	6.4	3.6
50282_at	hypothetical protein FLJ10975		AA522810	0	0.2	3.6
88783_at			AI961262	0.9	0.8	3.6
53745_at	LCHN protein		AI345949	0.2	0.7	3.6
74529_at			AI056998	0.4	0.2	3.6
53828_at			AI929168	5	1.1	3.6
59198_at	hypothetical protein		AF075050	7.2	4.3	3.6
52133_g_at			AI125264	0.3	1.9	3.6
80171_at			AI333006	7.2	1.8	3.6
62885_at			AI286024	1.8	0.8	3.6
63180_at			AI203344	1.5	2.1	3.6
51578_at			T91195	3.7	1.1	3.6
84140_at			AI623301	2.3	1.5	3.6
84900_at			AI219433	0.5	1.2	3.6
82705_at			AA464792	0.5	0.9	3.6
49200_at	serine/threonine kinase 24 (Ste20, yeast homolog)	STK24	AI364933	0	1.6	3.6
65361_i_at			HI6253	1.5	0.8	3.6
60052_at	hypothetical protein FLJ22662		AA534427	1.6	1	3.6
45363_at			AA161329	0	1.1	3.6
45233_at	mesenchymal stem cell protein DSC92		W22743	8.9	4.4	3.6
86740_at			AI458197	1.2	0.7	3.6
86376_at			AL042726	0.3	0	3.6
75830_at			AI791860	2.3	1.1	3.6
79035_at			H88339	10.5	6	3.6

Figure 13QQQQQQ

87660_at			AA700149	2.5	1.7	3.6
43555_at			AA282392	3.3	2.1	3.6
69266_at			AI634394	1.7	8.5	3.6
56275_at			AI631775	0.5	1	3.6
71407_at			AI651482	0.9	0.2	3.6
43503_at		GLS	AW003978	0.4	2.1	3.6
55506_at			AI828539	1.4	2	3.6
52675_at			AI581142	16.5	7.4	3.6
69003_at			AA602488	0.4	0.7	3.6
84811_at			AA992427	0.6	1.8	3.6
54806_at			AI807628	0	1.3	3.6
45347_at			AA533633	3.5	1.8	3.6
57198_at			AI636893	1.1	2.4	3.6
48531_at			AA522816	26	9	3.6
65202_at			AA651939	3.7	5.4	3.6
54887_at			AA772278	1.7	1.4	3.6
61663_at			AI825949	1.2	1.2	3.6
51220_at		SERF1A	W67895	13.1	4.7	3.6
60124_at			H61529	4.4	3.1	3.6
1390_s.at		GHRH	L00137	3	0.8	3.6
62110_at		growth hormone releasing hormone	N79004	2.6	0.7	3.6
53532_at			AI262173	1.5	1.3	3.6
51766_at			AI472196	0	0.2	3.6
46582_at			AA632424	0.8	1.7	3.6
46114_at			AA121481	3.7	0.6	3.6
46130_at			AL042817	0	0	3.6
60638_at			W74633	2.7	3.4	3.6
47384_at			AI806045	0	0.5	3.6
53757_at			AA131524	21.1	1.8	3.6
48742_at		activated RNA polymerase II transcription cofactor 4	AI979012	7.3	7.4	3.6
48520_at			AI638628	0.7	1.7	3.6
56675_at		epithelial membrane protein 1	H28004	2.3	2.5	3.6
75014_i.at		KIAA1533 protein	AW006583	64.4	37.7	3.6
42000_r.at			T40827	0.3	0.1	3.6
46050_r.at			AL120288	0.2	0.1	3.6

Figure 13RRRRRR

54324_at			AA206141	0.7	2.6	3.6
64811_at			AA234462	1.8	2.8	3.6
55558_at			AA524361	4.9	4.8	3.6
57766_r_at			AI391633	1.1	2.3	3.6
60895_at		SLC25A13	AA199865	1.8	1.3	3.6
57382_at			AI081773	2.6	1.2	3.6
59092_at			N50661	1.8	0.2	3.6
63467_f_at			AA084765	1.1	0.8	3.6
73314_at			W27255	3.7	1.4	3.6
88485_at			AI032311	1.2	0.6	3.6
86402_f_at			AA523697	1.4	0.2	3.6
40720_at			AL022398	2.8	0.5	3.6
39793_at		ADORA2BP	AF029786	1.8	1.3	3.6
32051_at		GBAS	AJ224875	3.5	1.8	3.6
36372_at		adenosine A2b receptor pseudogene				
33348_at		glioblastoma amplified sequence				
		hypothetical protein MGC2840 similar to a putative				
		glucosyltransferase				
		hexokinase 3 (white cell)	HK3	3.6	0.7	3.6
		transcription factor 12 (HTF4, helix-loop-helix transcription	TCF12	3.8	1.5	3.6
		factors 4)				
41309_g_at		C-terminal binding protein 1	U37408	3.2	2	3.6
38703_at		aspartyl aminopeptidase	AF005050	1.5	1	3.6
33916_at		imidazole receptor candidate	AB023192	4.4	3.8	3.6
37666_at		proteasome (prosome, macropain) subunit, beta type, 5	D29011	2.1	1	3.6
37555_at		PWP2 (periodic tryptophan protein, yeast) homolog	X95263	3.7	2	3.6
210_at		phospholipase C, beta 2	M95678	4.3	3.7	3.6
57346_at		Mad4 homolog	AA831438	2.5	3	3.6
59336_at		arrestin, beta 1	AI991333	3.6	2.6	3.6
57566_at		hypothetical protein	AA253487	3.1	3	3.6
56803_at		wingless-type MMTV integration site family, member 5A	AI968085	0	1.5	3.6
54862_at			W74437	2	2	3.6
49005_at		myosin regulatory light chain interacting protein	AI954822	3.8	3.4	3.6
68924_at			AA969377	3.7	3.1	3.6
65761_at			AA525969	3.9	2.8	3.6
44569_at			AA961420	3.8	3.6	3.6
64410_at		KIAA1150 protein	AA813014	4	2.6	3.6
52178_at			AA526595	1.8	2.1	3.6

Figure 13SSSSS

45690_at	CGI-133 protein		AA887118	2.7	1.9	3.6
80251_r_at	ATPase, Ca++ transporting, ubiquitous	ATP2A3	A1380736	3.6	3.6	3.6
57195_at			A1935271	3.7	2.2	3.6
57203_at	hypothetical protein FLJ10913		AA142864	2.5	3.1	3.6
46612_at	KIAA1624 protein		R39794	0.9	1.9	3.6
43399_s_at	KIAA0263 gene product		AW007492	3.5	2.6	3.6
47082_at	NAG-5 protein		AW003127	3.4	2.7	3.6
87754_g_at	serologically defined colon cancer antigen 3	SDCCAG3	W02630	3.6	3.3	3.6
63026_at	osteoclast stimulating factor 1	OSTF1	AA886976	1.1	1.3	3.6
46724_at	tensin	TNS	AL046979	0.4	0.8	3.5
41444_at	mitogen-activated protein kinase 7	MAP2K7	AF013588	7.3	2.1	3.5
37424_at	HCR (a-helix coiled-coil rod homologue)		AB029343	5.8	3.2	3.5
39361_f_at	transmembrane 4 superfamily member 6	TM4SF6	AF043906	0	0	3.5
39405_at	KIAA0266 gene product		D87455	1.9	2.5	3.5
41712_at	KIAA0560 gene product		AB011132	0	0.2	3.5
188_at	ephrin-B1	EFNB1	U09303	1.3	1.4	3.5
40202_at	basic transcription element binding protein 1	BTEB1	D31716	4.7	1.5	3.5
36411_s_at	ELAV (embryonic lethal, abnormal vision, Drosophila)-like 2	ELAVL2	U29943	1.3	0.4	3.5
35441_at	GLE1 (yeast homolog)-like, RNA export mediator	GLE1L	AF058922	0	1	3.5
33280_r_at	SA (rat hypertension-associated) homolog	SAH	X80062	0.5	0.1	3.5
32180_s_at	gene with multiple splice variants near HD locus on 4p16.3	EZH1	AB000461	1	0.9	3.5
32259_at	enhancer of zeste (Drosophila) homolog 1	PDE1B	AB002386	3.4	3.4	3.5
38921_at	phosphodiesterase 1B, calmodulin-dependent	H2BFS	U86078	4	2.7	3.5
32819_at	H2B histone family, member S	LCP2	AJ223352	4.3	1.2	3.5
39319_at	lymphocyte cytosolic protein 2 (SH2 domain-containing leukocyte protein of 76kD)		U20158	10.1	3	3.5
39344_at	transformer-2 alpha (htra-2 alpha)		U53209	3.4	1.1	3.5
34305_at	poly(rC)-binding protein 1	PCBP1	Z29505	0.7	0.6	3.5
39628_at	RAB9, member RAS oncogene family	RAB9	A1671547	12.8	7.4	3.5
1445_at	chemokine (C-C motif) receptor-like 2	CCR2	AF014958	32.3	6.4	3.5
772_at	v-crk avian sarcoma virus CT10 oncogene homolog	CRK	D10656	0	1.2	3.5
34892_at	tumor necrosis factor receptor superfamily, member 10b	TNFRSF10B	AF016266	5.4	0.3	3.5
34607_at	inducible T-cell co-stimulator	ICOS	AB023135	0.6	0.5	3.5
41828_at	methyl-CpG binding domain protein 1	MBD1	Y10746	1.2	0.4	3.5
35576_f_at	H2A histone family, member C	H2AFC	AL009179	5.8	0.9	3.5

Figure 13TTTTT

33987_at	ADP-ribosylation factor 1	ARF1	M36340	2.9	1.9	3.5
201_s_at	beta-2-microglobulin	B2M	S82297	3.4	1.5	3.5
36657_at	apolipoprotein C-II	APOC2	AA883870	0	1.6	3.5
40646_at	chemokine (C-X3-C) receptor 1	CX3CR1	U20350	0	0.8	3.5
38035_at	myotubularin related protein 6	MTMR6	AF072928	0.7	1.2	3.5
38439_at	nuclear factor (erythroid-derived 2) like 1	NFE2L1	L24123	12	7.5	3.5
32445_at	platelet-activating factor acetylhydrolase, isoform Ib, beta subunit (30kD)	PAFAH1B2	D63390	1.9	1.8	3.5
39137_at	nuclear factor related to kappa B binding protein	NFRKB	X80878	0.8	0	3.5
64955_at			A1435076	5.8	0.6	3.5
56134_at			A1458947	0.8	2	3.5
48475_at	hypothetical protein FLJ10748		R41648	9.2	0.9	3.5
57775_s_at			A1830218	1.7	2	3.5
64361_s_at	ubiquitin-conjugating enzyme E2L 3	UBE2L3	AW006924	0	0	3.5
50720_at	NAG14 protein		A1769160	0	1.5	3.5
65118_at	hypothetical protein FLJ10498		A1886468	3.4	2.1	3.5
56431_at	hematological and neurological expressed 1	HN1	N90863	0	0	3.5
75638_at			H88347	1.4	0	3.5
75814_at			A1218193	1.2	3.5	3.5
75843_at			AA808011	0.4	0.1	3.5
57446_at			AA687142	1.1	0.3	3.5
55908_at			A1239698	1.4	1.4	3.5
76587_at			AA868594	0.4	1.9	3.5
47199_at			AL119929	2.3	2	3.5
46529_at			AA988769	1.5	4.9	3.5
76411_at			A1798883	5.1	1.9	3.5
46214_at			AA521008	0	1.1	3.5
59169_at			A1733367	1	1.4	3.5
59910_at			AA908954	0.6	0	3.5
49924_at			A1074863	2.9	2.1	3.5
82355_at			A1358881	0.7	0	3.5
60670_at			A1096902	0.9	1.6	3.5
50503_at			AA203415	1.1	0.9	3.5
61324_at	cargo selection protein (mannose 6 phosphate receptor binding protein)		A1636693	3.8	3	3.5

Figure 13UUUUUU

91393_at		AA553560	1.8	1.3	3.5
81065_at		AI148152	2.6	1.5	3.5
81432_r_at		AI357605	2.4	0.7	3.5
84780_at		AI520664	1.6	1	3.5
47525_at	Kreisler (mouse) maf-related leucine zipper homolog	AI869550	0	3.7	3.5
46744_s_at	carboxypeptidase M	AA902480	5.3	3.6	3.5
62329_at		T51067	1.3	0.6	3.5
59507_at		AI638235	0	0.6	3.5
43068_at		T95617	0.3	1.3	3.5
49303_i_at	TBP-associated factor 172	AI989914	0	2.5	3.5
44851_s_at		N48088	8.8	8.4	3.5
87408_at		AI221305	4.6	0.3	3.5
68325_at		AI439633	3.2	1.4	3.5
68704_at		AI500310	0	1.2	3.5
49516_at		AA531016	5	6.6	3.5
44950_at		N92407	1	0.5	3.5
74036_at		AW015189	0.5	1.4	3.5
44069_at		H03287	2.7	6.1	3.5
50427_at		AW003857	10.5	7.2	3.5
49508_at		AI041540	9.1	5.2	3.5
68523_at		AA614814	6.6	2.6	3.5
81665_at		AI570315	1.8	1.5	3.5
82032_at		AI823999	4.6	5.2	3.5
45557_r_at	fibronectin 1	AW021977	3.6	1.4	3.5
45857_at		AI668561	0	4.2	3.5
53534_at		N63469	0.7	0.8	3.5
58564_s_at		N78066	3.4	3.2	3.5
53612_at		AI458702	1.4	0	3.5
33559_at	PTK6 protein tyrosine kinase 6	U61412	1.6	1	3.5
83721_at		W56091	0.7	0.9	3.5
64786_at		AI350962	0.2	2.2	3.5
51951_at		AW026036	0	1.8	3.5
63969_f_at	clusterin (complement lysis inhibitor, SP-40,40, sulfated glycoprotein 2, testosterone-repressed prostate message 2, apolipoprotein J)	AI982754	20.2	3	3.5

Figure 13VVVVVV

37022_at	proline arginine-rich end leucine-rich repeat protein	PRELP	U41344	5.8	1.8	3.5
47066_at	hypothetical protein FLJ20333		A1823905	0.6	1.4	3.5
85904_at			AA418669	0	1.5	3.5
60963_at			AA195088	2.4	1.3	3.5
47074_at	disrupter of silencing 10		A1125866	6.3	2.2	3.5
83128_at			A1536092	3.9	2.2	3.5
48728_r_at			W74754	0.3	0.1	3.5
42112_at			AA398813	0.1	0.1	3.5
42243_at			AA460394	1.9	2.1	3.5
64841_r_at			AA412155	2.5	4.1	3.5
50760_at			2837876	2.1	2.4	3.5
51592_at			2941536	1.8	1.6	3.5
52351_at			3053308	0.1	0.5	3.5
70054_i_at			T99829	4.8	1.7	3.5
1127_at	ribosomal protein S6 kinase, 90kD, polypeptide 1	RPS6KA1	L07597	2.6	3.5	3.5
40244_s_at	hypothetical protein FLJ11585		A1743654	3.4	2.6	3.5
33748_at	minor histocompatibility antigen HA.1		D86976	2.1	1.8	3.5
33908_at	calpain 1, (mu/I) large subunit	CAPN1	X04366	2.8	1.2	3.5
36482_s_at	ATPase, Ca++ transporting, ubiquitous	ATP2A3	Y15724	3.6	2	3.5
35338_at	paired basic amino acid cleaving enzyme (furin, membrane associated receptor protein)	PACE	X17094	2.8	3.1	3.5
36172_s_at	adaptor-related protein complex 3, delta 1 subunit	AP3D1	AF002163	1.9	1.1	3.5
52238_s_at	transforming, acidic coiled-coil containing protein 3	TACC3	A1990642	3.9	3.4	3.5
56336_at			H45858	2.7	3	3.5
48713_g_at	hypothetical protein FLJ11200		T84485	3.3	2.7	3.5
57917_at	HP1-BP74		N29669	2.3	1.9	3.5
57038_at	actin related protein 2/3 complex, subunit 4 (20 kD)	ARPC4	A1992284	2.2	1.6	3.5
60046_at	candidate tumor suppressor protein		A1796241	3.8	3.2	3.5
63258_at			A1499490	2.1	2.2	3.5
56238_at	PDZ-LIM protein mystique		A1589776	3.5	3.4	3.5
60791_at	hypothetical protein FLJ11011		A1828160	2.8	1.6	3.5
44047_at	hypothetical protein FLJ20160		AA133311	3.3	3.3	3.5
64036_at	hypothetical protein FLJ10716		A1968206	4.1	2.2	3.5
53614_at			A1160121	2.9	1.5	3.5
57165_at	hypothetical protein FLJ20287		AA088410	3.5	3.2	3.5

Figure 13WVWWW

53879_at	UDP-Gal:betaGlcNAc beta 1,3-galactosyltransferase, polypeptide 3	B3GALT3	AA167825	3.6	1.1	3.5
58963_at	hypothetical protein FLJ10535		W74642	4	2.4	3.5
65465_at			AI953614	3.6	2.8	3.5
45224_at			AA025862	5.1	3.2	3.5
57223_at	potassium voltage-gated channel, Isk-related family, member 3	KCNE3	AI982768	3.8	3.2	3.5
44143_at	McKusick-Kaufman syndrome	MKKS	AA399076	2.6	3.1	3.5
54547_at	hypothetical protein DKFZp434G0522		AA147848	3.5	2.2	3.5
57222_at			AA203532	3.9	2.6	3.5
54920_at	N-acetyltransferase, homolog of S. cerevisiae ARD1		AA075546	3.2	3.1	3.5
65978_at			AI817130	0	0	3.5
55653_at			AA524700	3.7	2.1	3.5
57299_s_at	related RAS viral (r-ras) oncogene homolog	RRAS	T54317	2.8	2.2	3.5
34435_at	aquaporin 9	AQP9	AB008775	50.5	15.3	3.4
39030_at	Rab acceptor 1 (prenylated)	RABAC1	AJ133534	1.6	3.1	3.4
39845_at	HtrA-like serine protease		AF020760	1.6	1	3.4
37189_at	phosphomannomutase 1	PMM1	AL023553	2.9	0.2	3.4
1158_s_at	calmodulin 3 (phosphorylase kinase, delta)	CALM3	J04046	0.4	0.1	3.4
39796_at	proteasome (prosome, macropain) activator subunit 3 (PA28 gamma; Ki)	PSME3	U11292	0.2	1.2	3.4
38566_at	collagen, type X, alpha 1 (Schmid metaphyseal chondrodysplasia)	COL10A1	X60382	1.3	0.9	3.4
1323_at	ubiquitin B	UBB	X04803	9.2	4.8	3.4
34195_at			AL121073	0.4	1	3.4
34678_at	fer-1 (C.elegans)-like 3 (myoferlin)	FER1L3	AL096713	3.3	8.2	3.4
315_at	zinc-finger DNA-binding protein		D45132	2.2	0.3	3.4
39131_at	SWI/SNF related, matrix associated, actin dependent regulator of chromatin, subfamily a, member 5	SMARCA5	N36842	0.5	1.5	3.4
34012_at	keratin, hair, acidic, 4	KRTHA4	Y16790	0.4	0	3.4
34948_at			AI820065	1	3.6	3.4
35845_at	SEC24 (S. cerevisiae) related gene family, member B	SEC24B	AJ131245	5.1	1.2	3.4
36953_at	MAD (mothers against decapentaplegic, Drosophila) homolog 4	MADH4	U44378	4.6	3.1	3.4
36980_at	proline-rich protein with nuclear targeting signal		U03105	14.4	2.2	3.4

Figure 13XXXXXX

36789_f_at	leukocyte immunoglobulin-like receptor, subfamily B (with TM and ITIM domains), member 5	LILRB5	AF025534	7.2	0.8	3.4
38122_at	solute carrier family 23 (nucleobase transporters), member 1	SLC23A1	D87075	0	0	3.4
58033_at			A1934556	3.2	0.4	3.4
57737_at			A1150441	3.2	5.1	3.4
64983_r_at			AA412585	4	1.5	3.4
58119_at			AA481066	3.6	1.9	3.4
56999_at			A1625959	28.4	11	3.4
56490_at			AA682764	0.4	0.5	3.4
76238_at			A1377797	3.3	2.6	3.4
59433_at			N32185	1.4	0.7	3.4
76396_at			AF150414	1.1	0	3.4
77406_at			A1806594	0.9	1.5	3.4
76536_at			A1693917	2.5	1.8	3.4
52492_at			W91997	2.2	1.3	3.4
79457_at			AW044435	0.2	1.7	3.4
61489_at			A1718763	12.9	3.7	3.4
77199_at			A1521904	2.1	1.7	3.4
80829_at			AA210905	5.2	4	3.4
50763_at	KIAA1080 protein; Golgi-associated, gamma-adaptin ear containing, ARF-binding protein 2		AA779863	5.1	6	3.4
58309_at			A1802060	2.6	4.7	3.4
82223_at			A1871160	3.9	2.3	3.4
84624_at			AA505694	0.6	0.5	3.4
62705_at			A1094787	2.4	0.2	3.4
44741_at			A1633622	0.6	1.2	3.4
45323_at	hypothetical protein FLJ20093		W87653	5.3	2.3	3.4
84750_at			AW005376	2.6	0.7	3.4
60437_at	PRKC, apoptosis, WT1, regulator	PAWR	AA527277	1.8	0	3.4
41747_s_at	MADS box transcription enhancer factor 2, polypeptide A (myocyte enhancer factor 2A)	MEF2A	U49020	0.8	1.5	3.4
67675_at			A1560389	4.5	0.7	3.4
43697_at			AA504249	7.2	3.3	3.4
51435_s_at			AA442383	1.3	0.4	3.4
79835_at			AA203646	0	0.1	3.4

Figure 13YYYYYY

59647_at		AA702942	2	1.1	3.4
42868_at		R61231	1	1.3	3.4
43025_r_at		AI821404	0.7	0.9	3.4
63908_r_at		R46086	2	4.9	3.4
43838_s_at	hypothetical protein PRO1068	AA100948	5.5	0.4	3.4
49980_at	hypothetical protein FLJ11078	AI201336	0	1.5	3.4
75736_at		AA625990	2.2	1.3	3.4
50081_at		AI765698	3.2	4.6	3.4
56288_at		AA190594	1.7	0.6	3.4
57923_r_at	KIAA1228 protein	R93394	1.4	1.3	3.4
43077_at		AA010208	0.3	2.3	3.4
70424_at		AI337065	1.4	0.1	3.4
44309_at		AI091158	1.8	1.9	3.4
65628_at		AW023171	27.4	1.6	3.4
62321_i_at		AA482950	3.4	2.6	3.4
47539_at		AI807478	0.5	0.6	3.4
44516_at		AI809013	1	0.2	3.4
41955_r_at		Z41561	2.6	0.9	3.4
44626_g_at		R78655	0.9	0.9	3.4
50965_at	RAB26, member RAS oncogene family	AI690165	2.8	1.7	3.4
45187_at		AW006235	7.3	0	3.4
46408_at		N21233	1.3	1.2	3.4
54352_at		AA210841	2.9	1.1	3.4
45785_at		AL040982	19.7	8.6	3.4
46900_at	DKFZP566B1346 protein	N62934	2.8	1.4	3.4
47331_at		W84346	2.6	2.4	3.4
53603_at		D79487	3.4	1.1	3.4
47797_at	nodal, mouse, homolog	AI050866	3	2.1	3.4
56861_at	RaP2 interacting protein 8	AI197870	1	0.4	3.4
45511_at		AI989557	1.5	0.9	3.4
45300_at	cyclic AMP phosphoprotein, 19 kD	AI525176	2.1	2.3	3.4
60546_at		N32556	1.5	0.5	3.4
42362_at	CCAAT/enhancer binding protein (C/EBP), delta	AI633245	13.4	3.1	3.4
72582_at	glutamate receptor, metabotropic 7	AI672542	0.2	1.8	3.4
48318_at		AA872062	3.8	1.9	3.4

Figure 13ZZZZZ

43417_r_at	CBF1 interacting corepressor		AA199649	2.8	3.4	3.4
47164_at			R56903	7.9	1.2	3.4
63060_i_at	cerebral cavernous malformations 1	CCM1	AA744866	1.6	3	3.4
55689_at			AW001579	0	2.2	3.4
47602_at			AA393223	5.7	0.1	3.4
38356_at	folistatin	FST	M19481	7.4	1	3.4
63296_at			AI741141	2.5	2.7	3.4
31535_i_at			W27858	0.4	0	3.4
56276_f_at			AA583394	1.6	1.1	3.4
51441_f_at			1924131	0.7	0.9	3.4
51480_at			2882851	2.6	2.3	3.4
70323_f_at			AI743616	3.5	4.4	3.4
33253_at	KIAA0129 gene product		D50919	2.2	1.5	3.4
35323_at	eukaryotic translation initiation factor 3, subunit 9 (eta, 116kD)	EIF3S9	U78525	3.3	1.9	3.4
37345_at	calumenin	CALU	AF013759	3	1.1	3.4
32610_at	LIM domain protein		X93510	3.4	2.4	3.4
479_at	disabled (Drosophila) homolog 2 (mitogen-responsive phosphoprotein)	DAB2	U53446	3.3	3	3.4
38270_at	poly (ADP-ribose) glycohydrolase	PARG	AF005043	3.6	1.3	3.4
38292_at	Homer, neuronal immediate early gene, 2		AF093264	2.9	0.5	3.4
58439_at	RP42 homolog		AI092548	1.1	1.3	3.4
58498_at	KIAA1378 protein		AI082236	3.4	2.5	3.4
58412_at	retinal degeneration B beta		AA807607	1.4	3.1	3.4
59298_at	hypothetical protein MPMGp800B12492Q3		AL079457	1.5	1.3	3.4
48498_at			AA868896	3	2.7	3.4
45132_at	hypothetical protein MDS025		AI341661	2.4	1.8	3.4
44794_s_at	replication factor C (activator 1) 1 (145kD)	RFC1	N37088	2.3	1.7	3.4
41222_at	signal transducer and activator of transcription 6, interleukin-4 induced	STAT6	AF067575	0.6	1.3	3.4
44850_s_at	hypothetical protein FLJ10024		AI985749	1.7	1.9	3.4
51173_at	purinergic receptor P2X, ligand-gated ion channel, 4	P2RX4	AI872092	3.4	2.9	3.4
57059_at			AA889576	2	2.1	3.4
57461_at			AI472111	3.5	3.2	3.4
64054_at	hypothetical protein PP3501		R36266	3.4	0.7	3.4
44684_at	vacuolar protein sorting 16 (yeast homolog)	VPS16	AI874233	2.2	2	3.4

Figure 13AAAAAA

64368_s_at	hypothetical protein FLJ10470		AW001184	3	2	3.4
45673_at	Gem-interacting protein		AA218963	1	0.7	3.4
46887_at			AI887726	2.8	0.9	3.4
46132_at			AA210738	3.8	2.1	3.4
53940_at			AI953658	3.3	3.1	3.4
54483_at			AA760777	3.4	3.4	3.4
65966_at	hypothetical protein		AA775776	2	1	3.4
48337_at			T59859	2.7	1.4	3.4
91390_at			AA508196	3.2	0.9	3.4
37150_at	Kelch motif containing protein		AB026190	1.2	0	3.3
39550_at	KIAA0584 protein		AB011156	0	0	3.3
39838_at	KIAA0622 protein; Drosophila "multiple asters" (Mast)-like homolog 1		AB014522	3.5	2.2	3.3
34903_at	KIAA1218 protein		AI017382	2.1	1.3	3.3
40166_at	CS box-containing WD protein		AA923149	0.2	1.1	3.3
33034_at	rhomboid (veinlet, Drosophila)-like	RHBDL	Y17108	2.7	1.3	3.3
40083_at	KIAA0625 protein		AB014525	0.1	1.7	3.3
32623_at	gamma-aminobutyric acid (GABA) B receptor, 1	GABBR1	AJ225028	1.7	1.6	3.3
40602_at			AF052178	3.4	2.2	3.3
2019_s_at	integrin, beta 7	ITGB7	M68892	0.3	1.6	3.3
40898_at	sequestosome 1	SQSTM1	U46751	2	4.8	3.3
32784_at	serine/threonine-protein kinase PRP4 homolog		AB011108	6.6	3.2	3.3
40446_at	PHD finger protein 1	PHF1	AL021366	3.3	2.8	3.3
35533_f.at	killer cell lectin-like receptor subfamily C, member 4	KLRC4	AJ001683	0	0	3.3
39603_at	disrupted in schizophrenia 1	DISC1	AB007926	2.8	0	3.3
37493_at			H04668	4.6	4.7	3.3
39076_s_at	DR1-associated protein 1 (negative cofactor 2 alpha)	DRAP1	AI991040	0.7	1.4	3.3
34873_at	nebullette		Y16241	0.3	0	3.3
33950_g.at	corticotropin releasing hormone receptor 2	CRHR2	AF011406	0.7	4.5	3.3
36131_at	dimethylarginine dimethylaminohydrolase 2	DDAH2	AJ012008	1.5	4.4	3.3
36196_at	phosphofructokinase, muscle	PFKM	U24183	2.3	2.5	3.3
37710_at	MADS box transcription enhancer factor 2, polypeptide C (myocyte enhancer factor 2C)	MEF2C	L08895	0.8	0	3.3
38070_at			AL080234	1.1	0	3.3
38071_at	heterogeneous nuclear ribonucleoprotein F	HNRPF	L28010	1.7	2.2	3.3

Figure 13BBBBBB

37637_at	regulator of G-protein signalling 3	RGS3	U27655	1.5	0	3.3
38470_i_at	amyloid beta precursor protein (cytoplasmic tail)-binding protein 2	APPBP2	D86981	0.7	0.3	3.3
37928_at	nuclear transcription factor Y, beta	NFYB	AA621555	0.3	0	3.3
37967_at	lymphocyte antigen 117	LY117	AF000424	0	1.3	3.3
38778_at	KIAA1046 protein		AB028969	5.3	4.8	3.3
33191_at	hypothetical protein FLJ10512		AW051579	0.8	0	3.3
31708_at	ribosomal protein L30	RPL30	L05095	1.3	1.3	3.3
42369_at			R43543	0	0.3	3.3
46759_at			R73588	1.8	0.3	3.3
42315_at			AA470110	3.4	1	3.3
76475_at			AI651996	1.1	2.1	3.3
59402_at			W58423	2.7	1.8	3.3
48821_at	small membrane protein 1		AI569248	1.1	2	3.3
66316_at			N24645	3.7	0	3.3
50840_r_at	neurexophilin 4	NXPH4	AI498771	5.6	2	3.3
76786_at			AI733079	3.1	0	3.3
73906_at			AW043793	0.9	2.5	3.3
76941_at			AI425072	1.5	1.1	3.3
57675_at			AW052119	7.8	5.1	3.3
53798_at	zinc-fingers and homeoboxes 1	ZHX1	AI123518	0.7	1.5	3.3
66893_f_at			AA904929	1.9	3.2	3.3
78943_at			AA909480	1.6	2.1	3.3
50541_at			AI732535	2	1.7	3.3
61217_at			AI090818	1.2	1.8	3.3
57053_s_at	homolog of rat nadrin		AI189504	0.1	0.8	3.3
79437_at			AI138554	0.4	0	3.3
52635_at			AI820880	0.1	0	3.3
62045_at	epidermal growth factor receptor substrate EPS15R		T05110	5.9	2.8	3.3
54636_at			R42627	1.1	1.3	3.3
83433_at			H30069	2.6	0	3.3
74978_at			AW025472	1	0.7	3.3
84239_at			R51644	1.5	1.3	3.3
91683_at			AI968374	3.8	2.3	3.3
44719_at	hypothetical protein FLJ10035		AA126848	2.7	2.2	3.3

Figure 13CCCCCCC

42787_at			AI141062	1.9	0.4	3.3
84162_at			AI457626	1.5	0	3.3
42818_r_at			R43093	2.2	2.9	3.3
85133_at			AI148712	3.9	1.3	3.3
84539_at		CRB1	AI805296	5.6	1.9	3.3
54789_at			AI376450	3.4	1.6	3.3
57525_at			AI738434	6.4	1.3	3.3
67058_at			AI204266	0.5	1.6	3.3
46309_at		TUBB	AI039239	3.4	1.3	3.3
54826_at			AA284268	7.8	4.8	3.3
88392_r_at			AI311557	1.1	0	3.3
59058_at			AA404351	0	0	3.3
87852_at			AI479665	0.7	0	3.3
43615_at			AA766300	1.9	2.5	3.3
68496_at			AI653006	4.3	1.3	3.3
49345_at			AI770110	1.1	6.7	3.3
70869_at			AI831416	1.6	2.1	3.3
71205_r_at			AI917887	2.3	0.8	3.3
72547_at			AI962788	2.4	2.5	3.3
43713_r_at			N57781	1.4	1.3	3.3
51185_at		PAN2 protein	AI309334	5.6	4.2	3.3
60939_i_at			AA151265	1.4	2.3	3.3
65554_at			AI188518	3.4	6.8	3.3
54578_at			AI123255	2.8	4.3	3.3
42365_f_at			H60126	10.8	6.9	3.3
62820_at			AW002871	0.4	0.6	3.3
45133_f_at			AI097467	2.6	2.4	3.3
61731_at			W69427	0.8	0	3.3
50978_at		hypothetical protein FLJ23342	AI338625	117.2	3.8	3.3
44945_at		putative secreted ligand homologous to ffx1	AA398841	0.3	2.2	3.3
46413_at		hypothetical protein FLJ20517	N22831	0.7	1.3	3.3
46419_at			AI826125	0.4	3.5	3.3
65080_r_at		RAB27A, member RAS oncogene family	AI557479	11.9	2.7	3.3
55262_at			W73113	1.5	2.4	3.3
47087_at			AI310524	6.2	2.5	3.3

Figure 13DDDDDDDD

48280_at		AA828461	7.9	1.7	3.3
50440_at		AA393461	0.9	0.6	3.3
57281_at		AI041170	2.4	2.2	3.3
14115_at		D26561	0.4	0.8	3.3
61252_f_at		H60379	0.6	0	3.3
52825_f_at		AI251752	0.8	0	3.3
67654_r_at		AA435672	0	0.6	3.3
72335_r_at		AI091278	1.9	1.1	3.3
92152_f_at		H08438	1.8	4	3.3
82498_at		AI479524	0	0.2	3.3
37845_at	hematopoietic protein 1	M58285	3.1	1.6	3.3
41278_at	BAF53	AF041474	3.8	3	3.3
34332_at	glucosamine-6-phosphate isomerase	D31766	3.6	0.9	3.3
40818_at	H-2K binding factor-2	D14041	3.8	2.4	3.3
39358_at	nuclear receptor co-repressor 2	U37146	2.1	0.9	3.3
36093_at	KIAA0614 protein	AB014514	3.3	3.1	3.3
40024_at	src homology three (SH3) and cysteine rich domain	D86640	3	3.1	3.3
32546_at	fumarate hydratase	U59309	2.3	2.1	3.3
1598_g_at	growth arrest-specific 6	L13720	2.2	2.6	3.3
38375_at	esterase D/formylglutathione hydrolase	AF112219	2.2	0.8	3.3
39160_at	pyruvate dehydrogenase (lipoamide) beta	D90086	3.5	2	3.3
39172_at		AL049319	3.5	2	3.3
48123_s_at		AI022930	3.1	2.1	3.3
90724_at		AA885833	3	3.3	3.3
57033_at	acetyl-CoA synthetase	N64822	2.8	2.4	3.3
80889_at		AA604308	3.7	2.5	3.3
54859_at	E2IG2 protein	AI609583	2.4	1.8	3.3
89945_s_at	filamin A, alpha (actin-binding protein-280)	AL048339	1.7	1.4	3.3
65916_i_at	hypothetical protein FLJ13660 similar to CDK5 activator-binding protein C53	AL045571	2.6	1.3	3.3
46743_s_at	myeloid/lymphoid or mixed-lineage leukemia (trithorax (Drosophila) homolog); translocated to, 7	AI814178	2.4	3.2	3.3
57132_at	RNB6	AI341482	3.5	3	3.3
43408_g_at	DKFZP586F1524 protein	AW014654	3.2	1.6	3.3
51101_f_at	tumor suppressor deleted in oral cancer-related 1	AI139045	1.1	1.3	3.3

Figure 13EEEEEE

64313_at	hypothetical protein FLJ20454		AW003067	2.9	3	3.3
64338_f_at	ribosomal protein S18	RPS18	AI990458	0.5	0	3.3
88830_s_at	EH domain-binding mitotic phosphoprotein		F30336	2.9	2.1	3.3
46623_at			AA579641	3.3	3.2	3.3
49106_at			AI972094	1.9	3.1	3.3
52550_s_at	hypothetical protein FLJ20727		AI633767	2	1.8	3.3
63999_s_at			AW008693	1.6	1.2	3.3
64584_at			H87175	3.1	3	3.3
62462_at	hypothetical protein dJ473B4		AI433432	3.5	3	3.3
43452_at			T15548	2.1	2.2	3.3
43072_s_at			AI690662	3	2.4	3.3
62552_at			AA287801	4.2	3	3.3
46262_at			AA532392	2.7	2.6	3.3
62559_at	hypothetical protein FLJ11230		W31010	3.4	1.6	3.3
75044_at	CGI-06 protein		AI968145	3.2	1.2	3.3
38657_s_at	clathrin, light polypeptide (Lca)	CLTA	M20471	0.4	1.3	3.2
41450_at	SAR1 protein		AI553878	2.4	0.7	3.2
39338_at	S100 calcium-binding protein A10 (annexin II ligand, calpactin I, light polypeptide (p11))	S100A10	AI201310	3.6	1.2	3.2
41654_at	adenosine deaminase	ADA	X02994	39.7	4.1	3.2
41864_at			AF052181	1.6	0.8	3.2
38137_at	FK506-binding protein 6 (36kD)	FKBP6	AF038847	3.9	0.5	3.2
31417_at	hypocretin (orexin) neuropeptide precursor	HCRT	AF041240	0	0.8	3.2
33598_r_at	chromosome 1 open reading frame 7	C1ORF7	AF054176	9.3	3.4	3.2
1294_at	ubiquitin-activating enzyme E1-like	UBE1L	L13852	2.5	5.8	3.2
39450_s_at	iduronate 2-sulfatase (Hunter syndrome)	IDS	AF050145	0.1	0.8	3.2
1170_at	colony stimulating factor 1 (macrophage)	CSF1	M37435	13.1	6.9	3.2
33261_at	major histocompatibility complex, class II, DR beta 1	HLA-DRB1	M16941	2.8	1.3	3.2
1548_s_at	interleukin 10	IL10	U16720	121.9	4.3	3.2
33794_g_at	transcription factor 20 (AR1)	TCF20	U19345	0.9	0.3	3.2
34215_at	DNA segment on chromosome X and Y (unique) 155 expressed sequence	DXYS155E	L03426	2.6	1.2	3.2
34292_at	chromosome X open reading frame 12	CXORF12	X92475	1.2	2.4	3.2
39568_g_at	aquaporin 7	AQP7	AB006190	1.1	1	3.2
33017_at	glycosylphosphatidylinositol specific phospholipase D1	GPLD1	AI040379	0.8	0	3.2

Figure 13FFFFFFF

178_f_at	postmeiotic segregation increased 2-like 8	PMS2L8	U38964	0.6	1.2	3.2
33368_at	protease, serine, 15	PRSS15	X76040	1.9	1.3	3.2
34778_at	NeuAc-alpha-2,3-Gal-beta-1,3-GalNAc-alpha-2, 6-sialyltransferase alpha2,6-sialyltransferase		AA418080	0.4	0.2	3.2
39548_at	neuronal PAS domain protein 2	NPAS2	U77970	1.7	0.8	3.2
34785_at	KIAA1025 protein		AB028948	3.9	2.3	3.2
34833_at	DKFZP586O0120 protein		AL050157	0	1.4	3.2
199_s_at	protein kinase C-like 2	PRKCL2	U33052	0.7	2	3.2
36671_at	asparagine synthetase	ASNS	M27396	16.8	9.8	3.2
36988_at	tumor necrosis factor, alpha-induced protein 1 (endothelial)	TNFAIP1	M80783	7.6	1.6	3.2
37544_at	nuclear factor, interleukin 3 regulated	NFIL3	X64318	8.9	8.7	3.2
33109_f_at	SRY (sex determining region Y)-box 2	SOX2	L07335	1	0.7	3.2
38822_at	serine/threonine kinase 17a (apoptosis-inducing)	STK17A	AB011420	3.2	1	3.2
41034_s_at	sulfoltransferase family, cytosolic, 2B, member 1	SULT2B1	U92315	0.6	0.9	3.2
56199_at	hypothetical protein FLJ20445		AI346493	3.2	2	3.2
56849_at			N99346	0.8	1.2	3.2
51174_at	spinocerebellar ataxia 7 (olivopontocerebellar atrophy with retinal degeneration)	SCA7	AL043992	3.9	4.4	3.2
59314_at	retinoic acid induced 1	RAI1	AI926671	5.7	4.5	3.2
76449_at			AA781217	8.8	0	3.2
59467_at	transmembrane phosphatase with tensin homology	TPTE	AA972714	13.6	4.7	3.2
90013_at			AA858297	0	1.1	3.2
77799_at			AI632915	1	0.7	3.2
76691_at			AI015524	1.3	0	3.2
50080_at			AI097194	5.2	3.3	3.2
91002_at			AI619500	2.2	0.9	3.2
61154_r_at			AA936133	0.6	2.5	3.2
78508_at			AI092265	2.1	0.4	3.2
62051_at			AI424912	2.4	1.6	3.2
83177_at			W63579	2.5	2.7	3.2
91400_at			AI309016	5.4	3.6	3.2
81534_at			AI422724	2	1.8	3.2
60036_at			AI125667	2.5	2.4	3.2
84063_at			AI417354	2.1	1	3.2
82297_at			AI435465	1.2	2.6	3.2

Figure 13GGGGGG

59364_s.at			R88719		2.3	1.9	3.2
82457_at			AI798172		0	0	3.2
43198_at			AI523598		1.3	1.1	3.2
85194_at			AI378091		1.5	3	3.2
67595_r.at			AI651263		0.8	1.8	3.2
58653_at	hypothetical protein from EUROIIMAGE 384293		AA952940		2	1.6	3.2
60479_s.at			AA779684		2	1.4	3.2
45258_at	hypothetical protein DKFZp761G2113		AF039240		9.2	4.7	3.2
43314_at			AA009521		1	1.6	3.2
86877_at			AA176617		4.3	0.4	3.2
51393_at			AA463643		2.8	1.7	3.2
63861_i.at	hypothetical protein FLJ10120		AI140364		1.5	0.7	3.2
88561_at			AI652609		1.9	2.4	3.2
79858_at			AI693951		6	5.7	3.2
59613_r.at			H78589		3.8	1.5	3.2
69998_at			AI764969		0.3	3.2	3.2
42902_at			W90753		0.4	1.1	3.2
71936_at			AI148006		5.6	4.2	3.2
44792_s.at			AA972102		0.8	5.2	3.2
50900_at			AI991581		0.5	0.9	3.2
45328_at	G protein-coupled receptor kinase 7	GPRK7	AI922109		1.4	3.3	3.2
59114_at			AA809355		1.9	1.8	3.2
63901_r.at			H55897		1.8	0	3.2
55526_at			AA923355		0.1	1.1	3.2
51237_at			AI754064		0	2.4	3.2
53128_at	hypothetical protein FLJ10494		AI678076		6.3	5	3.2
62643_at	hypothetical protein FLJ11029		AA292789		0	2.8	3.2
43895_i.at			F04022		2.8	2.1	3.2
63438_at			AI590098		2.7	0.4	3.2
73816_at			AW015805		1.7	0	3.2
83676_f.at			AI376179		2	1.1	3.2
58676_s.at	interleukin enhancer binding factor 1	ILF1	AI298446		0	1.5	3.2
61032_at			AI122634		0.3	0.4	3.2
48075_at	hypothetical protein FLJ20871 similar to FSP27		AI346596		2.6	2.6	3.2
44650_at			AI307750		6.2	2.7	3.2

Figure 13HHHHHHH

51706_at	hypothetical protein		AI816888	1.5	1.1	3.2
56915_at			AA595764	1.9	0.7	3.2
64242_at	peroxisomal biogenesis factor 11A	PEX11A	AI866665	2.6	2.2	3.2
44616_r_at			AA057585	0.5	0	3.2
58803_at	glucocorticoid modulatory element binding protein 1	GMEB1	AW003266	1	2.5	3.2
54379_at			N56947	3.8	0.4	3.2
64438_at	hypothetical protein FLJ22222		W19668	0.9	1.5	3.2
54083_at			AA167623	1.3	1.6	3.2
58646_at			AA824321	1.9	0.8	3.2
46297_at			AA149415	1.9	0.8	3.2
42055_at			AA282531	3.2	0.4	3.2
727_at				1	0.7	3.2
50735_at			2837330	0	1.3	3.2
68391_r_at			AI283981	0.6	1.3	3.2
67665_f_at			R02298	0.5	0	3.2
39507_at	O linked N-acetylglucosamine (GlcNAc) transferase (UDP-N-acetylglucosamine:polypeptide-N-acetylglucosaminyl transferase)	OGT	AL050366	1.3	1.2	3.2
38681_at	eukaryotic translation initiation factor 3, subunit 6 (48kD)	EIF3S6	U62962	1.5	1.3	3.2
34936_at	solute carrier family 4, sodium bicarbonate cotransporter, member 7	SLC4A7	AB012130	2.6	2.5	3.2
40452_at	copine 1	CPNE1	U83246	1.8	2	3.2
40571_at	myosin VA (heavy polypeptide 12, myosin)	MYO5A	U90942	3.1	0.7	3.2
32455_s_at	proline and glutamic acid rich nuclear protein		U88153	3.3	2.6	3.2
32506_at	KIAA1108 protein		AB029031	2.7	0.1	3.2
39637_at	solute carrier family 26 (sulfate transporter), member 2	SLC26A2	U14528	3.2	3.2	3.2
36001_at	methyltransferase-like 1	METTL1	Y18643	0.6	0.4	3.2
1924_at	cyclin H	CCNH	U11791	3.4	2.3	3.2
35266_at	bladder cancer associated protein	BLCAP	AL049288	3.7	1.8	3.2
40296_at	X-prolyl aminopeptidase (aminopeptidase P) 2, membrane-bound	XPNPEP2	AL023653	3.2	2.7	3.2
40745_at	adaptor related protein complex 1, beta 1 subunit	AP1B1	L13939	3.4	2.7	3.2
75779_at			AA629059	2.6	2.7	3.2
53779_at	CGI-147 protein		AI870849	1.7	1.4	3.2
57818_at	hypothetical protein dJ462023.2		AI363216	2.9	2.9	3.2

Figure 13IIIII

54865_at	coronin, actin-binding protein, 1C	CORO1C	N27805	3.2	2.9	3.2
54160_at	keratin 19	KRT19	A1056337	2.5	2.6	3.2
48577_at	src homology 3 domain-containing protein HIP-55		AW005562	1.2	0.9	3.2
63600_f_at			AW021173	3.1	2.9	3.2
60507_at			A1457453	0.7	0.9	3.2
45274_at	hypothetical protein		AA203167	3.3	1.8	3.2
49343_at			A1819630	2.9	2.9	3.2
58947_s_at	ribosomal protein L23	RPL23	AW020914	2.4	2.2	3.2
45562_f_at			AA913312	3.3	2	3.2
49242_at			N51263	3.2	2.6	3.2
43926_at			N36085	3.2	2.7	3.2
52691_at			AW014700	3.2	3.2	3.2
65816_at	hypothetical protein FLJ22378		AA210767	2.6	2.1	3.2
58523_at	HSPC182 protein		A1356771	2.1	1.8	3.2
44677_at	AD-015 protein		N42213	0.2	0	3.2
44696_at	hypothetical protein FLJ10743		AA915989	1.6	1.1	3.2
74729_f_at	anchor attachment protein 1 (Gaalp, yeast) homolog	GPAA1	AW007506	2.7	1.7	3.2
45795_at			A1672040	3.3	2.1	3.2
53159_at	hypothetical protein FLJ10829		A1056890	3.1	3.3	3.2
48511_at	TRIAD3 protein		H08320	2.8	1.4	3.2
55565_at	hypothetical protein FLJ20424		A1745491	3.2	2.5	3.2
48678_at	1,2-alpha-mannosidase IC		W22860	3.4	2.2	3.2
51695_s_at			AA649308	1.8	1.5	3.2
57723_at			AF063546	1.7	2.7	3.2
48910_at			A1742347	3	3.1	3.2
41235_at	activating transcription factor 4 (tax-responsive enhancer element B67)	ATF4	AL022312	6.7	4.1	3.1
41466_s_at	zinc finger protein 148 (pHZ-52)	ZNF148	L04282	0	0.2	3.1
39032_at	transforming growth factor beta-stimulated protein TSC-22		AJ222700	12.6	1.1	3.1
34480_at	cadherin 16, KSP-cadherin	CDH16	AF016272	2.6	2	3.1
37468_at	Janus kinase 2 (a protein tyrosine kinase)	JAK2	AF058925	3.7	4	3.1
39370_at			W28807	0.9	1.7	3.1
39742_at	TRAF family member-associated NFKB activator	TANK	U59863	5.2	4	3.1
40426_at	B-cell CLL/lymphoma 7B	BCL7B	X89985	2.8	0.6	3.1

Figure 13JJJJJ

40507_at	solute carrier family 2 (facilitated glucose transporter), member 1	SLC2A1	K03195	2.9	0.6	3.1
33227_at	interleukin 10 receptor, beta	IL10RB	A1984234	9.2	4.2	3.1
41221_at	phosphoglycerate mutase 1 (brain)	PGAM1	J04173	2.7	2	3.1
878_s_at	prolactin	PRL	M29386	0	0	3.1
32261_at	bromodomain adjacent to zinc finger domain, 1B	BAZ1B	AF072810	1.7	1.5	3.1
38899_s_at	hypothetical protein FLJ20693		U95822	4.3	5.7	3.1
38900_at	paired box gene 3 (Waardenburg syndrome 1)	PAX3	U02309	3.6	1.1	3.1
34089_at	KIAA1030 protein		AB028953	0.9	0	3.1
31655_at			AL031737	1	0.5	3.1
31699_at	phosphoinositide-3-kinase, catalytic, beta polypeptide	PIK3CB	S67334	0.4	0	3.1
35566_f_at	immunoglobulin heavy constant mu	IGHM	AF015128	1	0.9	3.1
34342_s_at	secreted phosphoprotein 1 (osteopontin, bone sialoprotein 1, early T-lymphocyte activation 1)	SPP1	AF052124	1.4	2.4	3.1
35802_at	KIAA1014 protein		AB023231	6.8	2.5	3.1
36618_g_at	inhibitor of DNA binding 1, dominant negative helix-loop-helix protein	ID1	X77956	1.2	0	3.1
38398_at	MAP-kinase activating death domain	MADD	AB002356	2.5	1.9	3.1
1794_at	cyclin D3	CCND3	M92287	0	0	3.1
38472_at	KIAA0143 protein		D63477	0.3	7.2	3.1
35030_i_at	KIAA0316 gene product		AB002314	1.2	0	3.1
38364_at	BCE-1 protein		AF068197	0	0.9	3.1
48788_at			AL043192	1.6	1.3	3.1
42389_at			AL039969	1.6	1.1	3.1
61118_at			AI983632	13.4	8.3	3.1
42445_at			AA608787	1.2	2	3.1
52939_at			N36115	3.7	1.7	3.1
74315_i_at	sorting nexin 16		AL039945	0.5	4	3.1
729_i_at	mucin 3A, intestinal	MUC3A	M55405	2	1.7	3.1
59882_f_at			AI150628	2	1.5	3.1
78690_at			AI939308	1.9	0.4	3.1
65433_at			T86971	0	0.1	3.1
51425_i_at			AA971774	3.6	1.5	3.1
79761_at			AI217928	2.1	4.5	3.1
61613_at			AI130697	1.1	2	3.1

Figure 13KKKKKKK

54611_s_at			W56695	0.8	1.6	3.1
80946_at			AA781852	1.1	1.2	3.1
83534_at			AI570127	1.2	0	3.1
62242_at			AA126691	2.6	2.4	3.1
84607_at			AA813111	1.3	0.5	3.1
82272_at			AI674070	2.1	1.6	3.1
82594_at			AI308862	5.8	6.1	3.1
63407_at	cell division cycle 27	CDC27	AA142869	0.9	0.8	3.1
83052_at			AA877124	4.1	2.9	3.1
51606_at			AI797595	1.9	0.1	3.1
83246_at			AI583921	0	0.3	3.1
86116_at			AI912018	1.2	0.4	3.1
48581_at			AA160530	1.2	0	3.1
45493_at	p30 DBC protein		N35906	0	1.5	3.1
67963_at			AA436007	4.7	4.7	3.1
67942_at			R08053	3.5	2.2	3.1
48231_at			H08161	0.8	0.8	3.1
88000_at			AA640065	0.8	1.3	3.1
68532_at			AA635985	2.2	0	3.1
42695_at			AA811286	2.8	0.8	3.1
69129_at			AI494189	1.1	2.1	3.1
86639_at			AI697562	0.3	0.5	3.1
49494_at			AI052684	1.9	1.7	3.1
50203_at	UDP-glucose:glycoprotein glucosyltransferase 2		AA883247	0	1.4	3.1
44758_at			R43528	1.9	1.2	3.1
44838_at	hypothetical protein FLJ12752		AI271348	1.3	4	3.1
52417_at			H04018	1.5	1	3.1
43553_at			AI807404	7.2	6.3	3.1
43869_at			R49539	0.9	0	3.1
45537_at	aquaporin 7	AQP7	F27435	2.2	3	3.1
49650_at			N68715	9.8	7.4	3.1
50710_i_at			H71301	2.7	2.3	3.1
65746_at	PC326 protein		N63302	0	1.1	3.1
56454_at	Breakpoint cluster region protein, uterine leiomyoma, 2		AI079838	3	1.7	3.1
90775_at			AI821755	0.9	0	3.1

Figure 13LLLLLLL

44026_at			AI741219	0.3	1.8	3.1
50318_at			AI497833	3.8	1.5	3.1
54491_at			AA043562	27.5	3	3.1
84545_at			AA601040	0.9	0.5	3.1
47969_at			AA631997	0	5.8	3.1
87221_at			AA829405	3.5	5.4	3.1
47328_at			AA483389	2.1	1.6	3.1
62139_at			AL043166	4.3	1.3	3.1
48440_at	hypothetical protein FLJ10874		AI767504	1.9	0.9	3.1
64247_at			H40631	20.6	4	3.1
50946_at			AI282277	1.8	1.4	3.1
771_s_at	CD7 antigen (p41)	CD7	D00749	3.1	0.9	3.1
64121_at	glioma-amplified sequence-41		Z78373	0	1	3.1
45073_at			N69578	0.1	1.7	3.1
47346_at			AI818064	0	0.3	3.1
54051_at			H59033	1.1	2.2	3.1
65107_at			AA028890	0.8	1.5	3.1
34640_at	transcription factor 1, hepatic; LF-B1, hepatic nuclear factor (HNF1), albumin proximal factor	TCF1	X71346	2.5	0	3.1
43436_g_at			AW023627	1.5	1.6	3.1
55007_at			AA521088	0.3	1	3.1
48617_at	disabled (Drosophila) homolog 1	DAB1	AI927931	1.1	1.4	3.1
65958_at	protein x 013		AA058963	1.5	0.3	3.1
48358_at			AA760689	0.9	1	3.1
46068_r_at			W70342	0.9	2.1	3.1
56060_at			AA913840	3.7	1.8	3.1
56072_at			AA939337	0.2	4.3	3.1
44986_s_at	DKFZP586N0721 protein		AI989895	0	2.8	3.1
1180_g_at			2837329	0	0.9	3.1
50734_at			AW007237	4	0.4	3.1
75712_at			AA586897	0.7	0	3.1
88334_f_at			AI620888	3.6	4.9	3.1
72603_at			U82938	2.6	1.7	3.1
39020_at	CD27-binding (Siva) protein		U01147	1.9	1	3.1
39058_at	active BCR-related gene	ABR				3.1

Figure 13MMMMMM

39910_at	hypothetical protein			AA663800	1.1	2	3.1
39783_at	KIAA0100 gene product			D43947	3.1	2.4	3.1
40045_g_at	chromosome 18 open reading frame 1		C18ORF1	AF009425	5.6	4.5	3.1
32060_at	BCL2/adenovirus E1B 19kD-interacting protein 2		BNIP2	U15173	2.1	0.5	3.1
40414_at	valyl-tRNA synthetase 2		VARS2	X59303	3.3	2.8	3.1
38242_at	B cell linker protein			AF068180	3.1	2.7	3.1
32825_at	HMT1 (hnRNP methyltransferase, <i>S. cerevisiae</i>)-like 2		HRMT1L2	Y10805	2	1.1	3.1
34661_at	KIAA0350 protein			AB002348	2.5	1.4	3.1
35145_at	MAX binding protein		MNT	X96401	2.8	1.5	3.1
34409_at	DKFZP564C1940 protein			AL080164	1.5	0	3.1
35714_at	pyridoxal (pyridoxine, vitamin B6) kinase		PDXK	U89606	2.2	0.1	3.1
35368_at	zinc finger protein 207		ZNF207	AF046001	1.2	0.8	3.1
35804_at	ash2 (absent, small, or homeotic, <i>Drosophila</i> , homolog)-like		ASH2L	AB022785	2.8	1.6	3.1
48488_at	hypothetical protein			AI889499	3.5	1.5	3.1
56521_at	hypothetical protein PRO1197			AI804075	1.3	1.1	3.1
57030_at				W26295	3.1	3.1	3.1
62889_at	KIAA0675 gene product			D29070	3.1	3.1	3.1
84776_at				AI817881	2.8	1.1	3.1
63713_at	serine/threonine kinase 24 (Ste20, yeast homolog)		STK24	AI524605	1.9	1.2	3.1
60471_at	hypothetical protein FLJ22439			AA625133	3.2	2.9	3.1
44083_at				AI479056	3.1	2	3.1
45251_g_at	KIAA1306 protein			AL039926	3	2.7	3.1
63323_at	hypothetical protein MGC3234			C05777	3	2.4	3.1
56293_at				AA150502	3.3	1.5	3.1
57091_at				T48293	2.2	1.9	3.1
43961_s_at	hypothetical GTP-binding protein DKFZp761H171			AI627648	2.3	0.9	3.1
44101_at				AI684986	3.1	2.7	3.1
43984_at				H15571	3.3	1.3	3.1
43992_at				R61748	3.6	3.3	3.1
47606_at	nucleoporin 50kD		NUP50	AI628114	2.9	2.3	3.1
44059_at	hypothetical protein PRO2605			AA151520	1.6	2.3	3.1
50400_at				AI743990	1.8	1.3	3.1
57160_at				N36348	2.1	1.9	3.1
52973_at	hypothetical protein FLJ10330			AA683501	3.5	2.6	3.1
50410_at				AI075950	0.9	0.2	3.1

Figure 13NNNNNN

45764_at	BUP protein		AI126303	3.4	2.7	3.1
51274_at			AA287396	1.8	1.9	3.1
74703_f_at	ATP synthase, H+ transporting, mitochondrial F0 complex, subunit c (subunit 9) isoform 3	ATP5G3	N91473	2.5	1.4	3.1
64377_at			AI925308	2.2	1.8	3.1
52169_at			AI302185	3.1	2.2	3.1
64420_at	hypothetical protein FLJ12085		AA873278	2.4	2	3.1
36500_at	NAD(P) dependent steroid dehydrogenase-like; H105e3		AF027974	2.9	1.5	3.1
61119_at	CCAAT/enhancer binding protein (C/EBP), alpha	CEBPA	AI971171	3.2	2.7	3.1
65299_at			C15819	3.1	3.3	3.1
37441_at	lipoyltransferase		AB017566	3.2	0.1	3
39899_at			AC005525	0.7	0.6	3
35012_at	myeloid cell nuclear differentiation antigen	MNDA	M81750	0	4	3
32049_f_at			AF070554	1.8	1.8	3
32073_at	KIAA0677 gene product		AB014577	4	1.3	3
709_at	tubulin, beta polypeptide	TUBB	J00314	0.4	0.5	3
32292_at	collectin sub-family member 10 (C-type lectin)	COLEC10	AB002631	0.6	0.3	3
40888_f_at	eukaryotic translation elongation factor 1 alpha 1	EEF1A1	W28170	1.3	1	3
40949_at	dynamin, cytoplasmic, light intermediate polypeptide 2	DNCL12	AF035812	0.4	2.3	3
33741_at	CGI-11 protein		AI741756	0.8	6.5	3
448_s_at	multiple endocrine neoplasia I	MEN1	U93237	1.7	0.7	3
35996_at	ZW10 interactor anti-sense	ZWINTAS	X98261	1.8	1.1	3
37488_at	farnesyltransferase, CAAX box, beta	FNTB	L00635	0.7	0.4	3
1046_at	retinoic acid- and interferon-inducible protein (58kD)		U34605	7	5.4	3
38085_at	chromobox homolog 3 (Drosophila HP1 gamma)	CBX3	AI740522	2	1.5	3
41491_s_at	ATPase, Class VI, type 11A	ATP11A	AB028944	3.3	0	3
35083_at	ring finger protein 24	RNF24	AL031670	2.1	1.2	3
35695_at	Chediak-Higashi syndrome 1	CHS1	U67615	2.5	2.2	3
34396_at	KIAA0978 protein		AB023195	3.2	2	3
34796_at	translocating chain-associating membrane protein		X63679	0	0.6	3
36726_at	paired box gene 1	PAX1	AL035562	1.5	0	3
35767_at	ganglioside expression factor 2		AI565760	1.3	2.2	3
37402_at	ribonuclease, RNase A family, 1 (pancreatic)	RNASE1	D26129	0.6	0.3	3
40403_at	myocilin, trabecular meshwork inducible glucocorticoid response	MYOC	Z97171	1	1.3	3

Figure 13000000

38287_at	proteasome (prosome, macropain) subunit, beta type, 9 (large multifunctional protease 2)	PSMB9	AA808961	15.2	15.8	3
39136_at	oxidative-stress responsive 1	OSR1	AB017642	5.7	3.1	3
38343_at	KIAA0328 protein		AB002326	0.8	1	3
58055_at			AI023433	0.5	1	3
48052_at			AI936906	0.5	1.9	3
42419_at			R42241	1.2	1.3	3
39007_at	matrix metalloproteinase 2 (gelatinase A, 72kD gelatinase, 72kD type IV collagenase)	MMP2	M55593	2.9	0.3	3
75980_at			H71242	1.3	2.5	3
76699_at			AA857941	1.7	2.6	3
63487_r_at			N71157	2.5	1.5	3
91017_at	baculoviral IAP repeat-containing 3	BIRC3	U54711	19.2	19.6	3
59873_at			AI217670	2.2	1.9	3
82144_at			AI917494	1.9	1.2	3
78077_at			AI125204	14.5	6	3
42555_at			T82467	1.4	0.4	3
60328_at			AI043376	0.6	1.4	3
74867_at			AI038997	3.7	4.3	3
90940_at			AI633529	0	1.8	3
52013_at			AI017881	1.4	1.2	3
79999_at			AI733807	1.3	0.8	3
83036_at			AI271559	5	0.5	3
80559_at			AI808359	2.9	2.8	3
54744_at			AA582287	23.5	8	3
81125_at			AI221390	1.5	4.3	3
83716_r_at			AI380017	0	1.5	3
84004_at			AI141609	1.8	0.9	3
84587_r_at			AA447648	2.3	1	3
65077_at	hypothetical protein FLJ13163		AW009360	2.3	1.8	3
90732_at	splicing factor, arginine/serine-rich 5	SFRS5	W45137	2.1	8.1	3
73674_r_at	glutamate receptor, metabotropic 4	GRM4	AI671792	1.9	1.1	3
55909_at			AI805301	0.2	0.3	3
48546_s_at	eukaryotic translation elongation factor 1 alpha 1	EEF1A1	N30008	0.6	1	3
63276_r_at			H53118	2.4	1.1	3

Figure 13PPPPPP

67647_at		AI479947	1.2	0	3
55932_at		R02328	1.4	1.6	3
68050_at		AA506364	0.6	1.5	3
58742_r_at		AA401418	0.9	1.6	3
88795_at		AI684457	0.8	1.3	3
69051_at		AI688514	2.3	1.4	3
89133_at		AA983457	1.5	0.7	3
69631_r_at		AI760376	0	0.9	3
70277_at		AI792561	1.9	1.8	3
56295_at		AI310158	2.6	2.2	3
71928_at		AI985592	3	1.3	3
49615_g_at	chromosome 20open reading frame 3	AA057573	2	1.7	3
55888_r_at	ALL1 fused gene from 5q31	AA704245	2.4	0.3	3
52685_at		AI991451	3.1	1.5	3
64755_at	hypothetical protein FLJ20758	AI147603	1	2.2	3
45680_at	KIAA1474 protein	AI741882	0	0	3
75929_at	muscleblind (Drosophila)-like	AI697540	2.8	0.2	3
46947_at		T87245	2.1	1.2	3
47273_at		AA252348	6.9	1.2	3
68818_at		AI198919	1.5	1.3	3
56245_r_at		AI469796	0.6	0.4	3
57074_s_at		AW021642	5.4	1.5	3
68083_at	KIAA0214 gene product	AA761811	7.6	2.1	3
45155_at	claudin 15	AI433892	1	0.9	3
46376_at		W73195	0.4	1.5	3
56776_at		W95146	1.8	1.1	3
92097_at		AI935368	0.2	0.5	3
51848_at		AI685919	1.6	0.8	3
58764_at	suppression of tumorigenicity 7	AI821369	2.2	3.7	3
58339_at	hypothetical protein FLJ13222	R52747	5	2.7	3
53970_at	dual adaptor of phosphotyrosine and 3-phosphoinositides	AF150266	43.8	9.5	3
38458_at	cytochrome b-5	L39945	0.5	2.7	3
63009_at	hypothetical protein FLJ10539	AI188402	0.1	0	3
56709_at		AI826531	1.6	0.1	3
44185_at	programmed cell death 6-interacting protein	AA156791	1.2	2	3

Figure 13QQQQQQ

54456_at			AL044991	2.1	1.1	3
61407_at			AA470003	3.4	2.8	3
56033_at	similar to rab11-binding protein		AI016239	0.5	2.2	3
58388_at			AI394215	1.6	2.4	3
52892_at			AI761092	3	0.5	3
70409_at			AI792564	3	0.8	3
72751_at			N67652	0.7	0.3	3
88372_at			AA708240	1.1	0.1	3
89249_at			T97523	0	0.1	3
67220_at			AW005511	5.3	3	3
1933_g_at	ATP-binding cassette, sub-family C (CFTR/MRP), member 5	ABCC5	U83661	3.1	2.5	3
38976_at	coronin, actin-binding protein, 1A	CORO1A	D44497	3.1	3.1	3
39864_at	cold inducible RNA-binding protein	CIRBP	D78134	1.1	0.3	3
33816_at	myosin IXB	MYO9B	AF020267	1.1	0.4	3
41758_at	chromosome 22 open reading frame 5	C22ORF5	AL096879	3.7	2	3
32181_at	flotillin 2	FLOT2	M60922	2.3	2.9	3
34291_at	phenylalanine-tRNA synthetase-like	FARSL	U07424	3.2	1.6	3
39424_at	tumor necrosis factor receptor superfamily, member 14	TNFRSF14	U70321	0.7	0.2	3
35723_at	nudix (nucleoside diphosphate linked moiety X)-type motif 1	NUDT1	D16581	4	3.1	3
36312_at	serine (or cysteine) proteinase inhibitor, clade B (ovalbumin), member 8	SERPINF8	L40377	0	0	3
41600_at	proliferation-associated 2G4, 38kD	PA2G4	U59435	3.1	2.9	3
35814_at	dendritic cell protein		AF064603	0.8	1	3
1138_at	solute carrier family 20 (phosphate transporter), member 1	SLC20A1	L20859	2.8	1.8	3
38481_at	replication protein A1 (70kD)	RPA1	M63488	3.3	2.3	3
38000_at	pyruvate carboxylase	PC	S72370	3.1	2.6	3
57931_s_at	6-phosphogluconolactonase	PGLS	AA582687	2	2.1	3
48053_at	hypothetical protein FLJ10416 similar to constitutive photomorphogenic protein 1		AI126869	2.6	2	3
65125_at	dolichyl-phosphate mannosyltransferase polypeptide 3	DPM3	AI817330	3	2.7	3
57005_at	histone deacetylase 8		AI972846	2.8	2.1	3
47431_at			AI742206	3.6	1.5	3
53730_at			W60775	3	2.9	3
77028_at			AA919126	2.6	0.5	3

Figure 13RRRRRR

58579_s_at	p47			W07096	3.2	2.3	3
57809_at	presenilins associated rhomboid-like protein			W63675	2.5	2.8	3
58858_s_at	CGI-07 protein			AW004635	2.6	1.7	3
47994_at	KIAA0788 protein			AI183987	2.9	1.3	3
67319_at				AI911560	2.2	2	3
60495_at				AI858726	2.9	1.9	3
60866_s_at	NS1-binding protein			AA535109	1.9	1.2	3
49463_at	suppressor of fused			AI760042	3.3	3.1	3
73349_at				AI949213	2	1.3	3
65612_at	hypothetical protein MGC3180			AW006145	3	3	3
62965_at	protein tyrosine phosphatase, non-receptor type 23		PTPN23	C75369	2.2	2	3
45542_f_at	mitochondrial import receptor Tom22			AI359916	2.4	2.1	3
49624_at				AI768630	2.2	0.7	3
54625_at	solute carrier family 26, member 6		SLC26A6	AA057543	2.4	2.1	3
64212_at				AI125474	2.8	2.4	3
64096_at	WD repeat domain 3		WDR3	AI880771	2.9	2.1	3
51230_at				AA293641	1.9	1.5	3
52094_at	similar to glucosamine-6-sulfatases			AI688582	3	2.8	3
52124_at				AI805463	3	2	3
62460_f_at				AA604243	2.9	2.4	3
46226_at	HSPC142 protein			AA610145	2.6	2.3	3
64517_at	KIAA0747 protein			AI972594	3.2	2.4	3
63022_at	hypothetical protein MGC3162			AI675158	2.8	2.4	3
59666_at	complement component C1q receptor			W72082	2.6	2.8	3
41108_at	Pseudoautosomal GTP-binding protein-like			Y14391	2	0.5	2.9
33815_at	uridine monophosphate synthetase (orotate phosphoribosyl transferase and orotidine-5'-decarboxylase)		UMPS	J03626	2.7	2.2	2.9
41243_at	KIAA0447 gene product			AB007916	2.9	2	2.9
34680_s_at	KIAA0107 gene product			D14663	1.4	0.8	2.9
35167_at	KIAA0433 protein			AB007893	3	2.1	2.9
34837_at	KIAA0376 protein			AB002374	3	1.9	2.9
35796_at	protein tyrosine kinase 9-like (A6-related protein)		PTK9L	Y17169	3	1.4	2.9
36135_at	nucleolar protein p40; homolog of yeast EBNA1-binding protein			U86602	2.9	2.7	2.9
37047_at	Niemann-Pick disease, type C1		NPC1	AF002020	1.5	0	2.9

Figure 13SSSSSS

36792_at	tropomyosin 1 (alpha)	TPM1	Z24727	3	3	2.9
37281_at	KIAA0233 gene product		D87071	1.7	1.6	2.9
38336_at	KIAA1013 protein		AB023230	3	2.5	2.9
57539_at			AA535065	0.8	1	2.9
48271_at			AI683752	2.9	2.8	2.9
78442_at	hypothetical protein MGC4692		AI819340	2.9	2.9	2.9
47463_at	eukaryotic translation initiation factor 2C, 1	EIF2C1	AA453723	3.2	3.1	2.9
58323_at			AA703241	2.8	0.8	2.9
74694_s_at	hypothetical protein FLJ23282		AA907940	1.5	0.9	2.9
51614_at			N98756	2.7	2.7	2.9
51034_at			AI761602	2.9	2.7	2.9
65750_f_at	hypothetical protein		N75645	3.1	0.1	2.9
56274_at	KIAA1303 protein		H06513	2.1	2.6	2.9
74544_at			AI769930	1.9	3	2.9
63889_at	KIAA1238 protein		AI760919	2.8	1.7	2.9
61400_i_at			AI952593	0	0.7	2.9
89584_at	protease, serine, 15	PRSS15	AI817597	2.5	0.9	2.9
48527_at			AI394438	1.7	1.5	2.9
59452_at			AI719167	2.8	2.8	2.9
43854_at			AW004730	3	2.8	2.9
50979_s_at			AI968170	2.9	2.1	2.9
46403_at			AI984021	3	2.6	2.9
44849_at	beta-site APP-cleaving enzyme	BACE	AI356651	2.9	2.9	2.9
46871_r_at			W32096	3.4	3.1	2.9
42054_at	signal transducer and activator of transcription 2, 113kD	STAT2	AA281793	2.2	2.5	2.9
54956_at			AA045241	2.7	1.2	2.9
90001_s_at	casein kinase 1, epsilon	CSNK1E	AI831641	3.3	2.8	2.9
47611_at	phosphodiesterase 4A, cAMP-specific (dunce (Drosophila)-homolog phosphodiesterase E2)	PDE4A	AA523395	2.6	2.1	2.9
47156_at			AI032786	3.2	2.5	2.9
57349_at			AA418743	2.5	2.7	2.9
55735_at			W61056	2.9	1.6	2.9
39812_at	mitochondrial ribosomal protein L12	MRPL12	X79865	2.7	2.2	2.8
40801_at	DKFZP434C212 protein		AA643063	3.1	1.3	2.8
39437_at			Z78324	2.4	1.5	2.8

Figure 13TTTTTTT

33838_at	HLA-B associated transcript-3		M35519	1.5	0	2.8
36935_at	RAS p21 protein activator (GTPase activating protein) 1	RASA1	M23379	3.1	1.8	2.8
37652_at	calcineurin binding protein 1		AB002328	2.7	2.1	2.8
37723_at	cyclin G2	CCNG2	U47414	2.9	1.6	2.8
38732_at	chloride channel, nucleotide-sensitive, 1A	CLNS1A	X91788	1.4	1	2.8
38250_at	KIAA0007 protein		D26488	2.9	2.7	2.8
48852_at	hypothetical protein DKFZp434H0820		AI346327	2.5	1.8	2.8
59680_at	hypothetical protein FLJ20548		AA523383	1.7	2.7	2.8
57340_s_at	skb1 (S. pombe) homolog	SKB1	AI969686	1.5	1.9	2.8
52271_s_at	peptide deformylase-like protein		AA516472	2.7	1.7	2.8
65710_at	NS1-associated protein 1		AI625087	2.4	1.9	2.8
44712_at			AI139894	3.8	1.8	2.8
82998_at			AI380429	3	2.6	2.8
86192_at			AA913255	3.1	2.7	2.8
48595_at	hypothetical protein FLJ10210		AA707400	3	2.8	2.8
59091_at			AI799698	2.5	2.4	2.8
45475_f_at	hypothetical protein FLJ10120		AW006786	2.5	1.3	2.8
69646_i_at			AA758581	2.8	2.5	2.8
71094_at			AA027791	2.8	2.2	2.8
50219_at	mRNA for FLJ00023 protein		W27853	3	1.3	2.8
43542_at			W75954	2.9	2.1	2.8
44740_at	hypothetical protein FLJ20343		W73280	2.6	1.8	2.8
73587_at	taln	TLN	AI494397	2.5	2.2	2.8
59771_at			T49250	2	1.7	2.8
47559_at			N66622	2.3	2.6	2.8
48729_at			AI126706	2.3	0.3	2.8
55498_r_at			AA171850	1.7	0.8	2.8
57918_at	DKFZP586G1517 protein		AA779617	2.8	2.6	2.8
52143_at	hypothetical protein FLJ12479		AI762566	2.3	1.4	2.8
45817_at	hypothetical protein		W31092	2.1	2.5	2.8
53956_at			AA449296	3.2	2.6	2.8
48747_at	phosphatidylinositol-4-phosphate 5-kinase, type II, beta	PIP5K2B	AI982873	2.5	2.3	2.8
46230_at			W80987	1.8	2.2	2.8
62548_at			AI655782	2.4	1.7	2.8
46735_s_at			AI952097	1.7	2.3	2.8

Figure 13UUUUUU

74537_f_at	alpha-2-macroglobulin	A2M	AI969632	2.4	0.7	2.8
46263_at			AA883486	2	0.4	2.8
74557_s_at	seladin-1		AI739473	2.8	2.6	2.8
46279_at			AA127629	2.9	2	2.8
46281_at	zinc/iron regulated transporter-like	ZIRT1	W21970	2.6	2.4	2.8
61289_at	ATPase, Ca++ transporting, plasma membrane 1	ATP2B1	AI337321	2.5	2.5	2.8
55091_at	phosphoinositol 3-phosphate-binding protein-2		AI341312	2.6	2.5	2.8
47139_at			AA481256	2.5	2.1	2.8
55671_at	FH1/FH2 domain-containing protein		N90883	2.9	2.7	2.8
54507_at			AA912049	3.4	1.9	2.8
37510_at	syntaxin 8	STX8	AF036715	2.3	1.6	2.7
39431_at	aminopeptidase puromycin sensitive	NPEPPS	AI132583	2.9	1.7	2.7
40207_g_at	chymotrypsin-like	CTRL	AI767675	1.8	1.6	2.7
38499_s_at	myelin-associated oligodendrocyte basic protein	MOBP	D28113	2.7	2.7	2.7
40817_at	nucleobindin 1	NUCB1	M96824	1.2	0.1	2.7
31873_at	Rho GTPase activating protein 4	ARHGAP4	U52112	4	2.8	2.7
41220_at	MLL septin-like fusion	MSF	AB023208	1.5	1.8	2.7
1468_at	heat shock protein 75		U12595	3	1.6	2.7
32212_at	programmed cell death 8 (apoptosis-inducing factor)	PDCD8	AL049703	2.7	2	2.7
33458_r_at	H2B histone family, member L	H2BFL	AI688098	2	1.4	2.7
39615_at	KIAA1026 protein		AB028949	2.8	2.6	2.7
39516_at	hypothetical protein		AI827793	2.9	2.2	2.7
34386_at	methyl-CpG binding domain protein 4	MBD4	AF072250	2.8	1.8	2.7
36028_at	T-cell, immune regulator 1	TCIRG1	U45285	1.4	0.6	2.7
36030_at	DKFZP586I2223 protein		AL080214	2.9	1.5	2.7
36732_at			AI004207	2.1	1.2	2.7
36164_at	Pyruvate dehydrogenase complex, lipoyl-containing component X; E3 binding protein		U82328	4	2.6	2.7
371_at	interferon regulatory factor 3	IRF3	Z56281	2.9	1.4	2.7
38021_at	plectin 1, intermediate filament binding protein, 500kD	PLEC1	U53204	2.7	1.5	2.7
38370_at			U90902	2.1	1.2	2.7
37647_at	acyloxacyl hydrolase (neutrophil)	AOAH	M62840	3	2.2	2.7
38770_at	a disintegrin and metalloproteinase domain 12 (meltrin alpha)	ADAM12	AF023476	2.9	2.9	2.7

Figure 13VVVVVV

38257_at	NADH dehydrogenase (ubiquinone) Fe-S protein 8 (23kD) (NADH-coenzyme Q reductase)	NDUFS8	AF038406	2.6	1.7	2.7
38290_at	regulator of G-protein signalling 14	RGS14	AF037195	2.9	2.6	2.7
38306_at	brefeldin A-inhibited guanine nucleotide-exchange protein 1		AA477576	2	1.1	2.7
33451_s_at	ribosomal protein L22	RPL22	AI526079	2.6	2.1	2.7
53721_at			AI742931	2.6	2.5	2.7
49349_at	PP2135 protein		W72919	2.6	2.6	2.7
52583_s_at	BCL2/adenovirus E1B 19kD-interacting protein 3-like	BNIP3L	AW025487	1.6	2.1	2.7
49204_at	glyoxylate reductase/hydroxypyruvate reductase	GRHPR	AI983536	2.1	1.6	2.7
73251_f_at	KIAA0741 gene product		AI990264	1.9	1	2.7
75220_s_at	filamin A, alpha (actin-binding protein-280)	FLNA	AW005498	1.4	0.9	2.7
65519_at	HLA class II region expressed gene KE2	HKE2	AI476588	1.9	1.8	2.7
49446_at			AI655698	2.9	1.9	2.7
57112_at	dipeptidylpeptidase III	DPP3	AI808417	1.9	2.3	2.7
50396_at	chromosome 12 open reading frame 5	C12ORF5	AI979251	2.2	2.3	2.7
43401_at			AI742497	2.6	1.3	2.7
56342_at	nuclear prelamin A recognition factor		AI963118	2.2	2.2	2.7
43977_at	hypothetical protein FLJ20422		AI660497	2.6	2.5	2.7
54927_at	eukaryotic translation initiation factor 2B, subunit 3 (gamma, 58kD)	EIF2B3	AI937500	2.7	2.5	2.7
57862_at			AA460579	2.3	2.5	2.7
60079_s_at	hypothetical protein FLJ20920		AA999701	2.9	2.4	2.7
88959_at			AA528441	2.5	1.5	2.7
63808_at			H23562	2	1.2	2.7
45114_at			AI815763	2.8	2.5	2.7
51788_at	proteasome (prosome, macropain) 26S subunit, non-ATPase, 1	PSMD1	AW023096	2.1	1.6	2.7
51085_at			AW005054	2.6	1.8	2.7
53015_at	hypothetical protein FLJ20729		AI823408	2.7	2.2	2.7
53061_at	glycoprotein M6B	GPM6B	AI718385	1.6	1.8	2.7
80466_s_at	hypothetical protein FLJ23439		AI394307	2.5	1.8	2.7
46677_at			AI762760	3	1.6	2.7
45360_s_at	dishevelled 1 (homologous to Drosophila dsh)	DVL1	AI990838	2	1.1	2.7
64383_at	tryptophanyl-tRNA synthetase	WARS	N66434	2	2.3	2.7
86462_at	bromodomain adjacent to zinc finger domain, 2B	BAZ2B	AI674297	2	1	2.7

Figure 13 WWWWWW

41696_at				AI620381	1.9	1	2.6
35172_at	tyrosylprotein sulfotransferase 2	TPST2		AF049891	2.5	2.2	2.6
35326_at	putative transmembrane protein; homolog of yeast Golgi membrane protein Yip1p (Yip1p-interacting factor)			AF004876	3.3	1.6	2.6
35342_at				AF052159	2.9	2.5	2.6
36978_at	KIAA0077 protein			D38521	1.5	0.4	2.6
37355_at	steroidogenic acute regulatory protein related			D38255	2.3	0.9	2.6
40191_s_at	KIAA0582 protein			AI761647	2.5	1	2.6
37597_s_at	similar to <i>S. cerevisiae</i> Sec6p and <i>R. norvegicus</i> rsec6			AF055006	2.7	2	2.6
38841_at	putative glioblastoma cell differentiation-related			AF068195	2.9	2	2.6
55748_at	ribosomal protein L17 isolog			AI141700	2.3	1.7	2.6
56197_at	phospholipid scramblase 3			AI783924	2.5	2.3	2.6
49141_at	GK002 protein; glib protein			AA631191	2.3	0.9	2.6
56953_at	uncharacterized hematopoietic stem/progenitor cells protein MDS031			AA126799	2.4	1.7	2.6
65126_at	major histocompatibility complex, class II, DN alpha	HLA-DNA		AI692538	2.6	2	2.6
56533_at	sorting nexin 8			N38744	1.9	1.6	2.6
44514_at	hypothetical protein FLJ12934			AA126458	3	1	2.6
53785_at	hypothetical protein FLJ10044			AA522674	2.3	2.4	2.6
80157_at				AI130690	2.7	2.1	2.6
71534_at				AA584378	1.3	1.2	2.6
44714_at	putative G-protein coupled receptor			AI261382	2.5	1.1	2.6
49686_s_at	vesicle-associated membrane protein 8 (endobrevin)	VAMP8		AI087923	2.5	1.5	2.6
85006_at				AA601045	2.6	2.6	2.6
60493_at				AI344328	2.7	2.5	2.6
32800_at	retinoid X receptor, alpha	RXRA		U66306	3	2.9	2.6
59644_at	hypothetical protein DKFZp434P0116			AI735391	0.6	2.3	2.6
78002_at				AI760531	2.6	1.9	2.6
43447_at	serologically defined colon cancer antigen 10	SDCCAG10		AI768148	2.6	1.2	2.6
43834_s_at				N68645	2.1	1.8	2.6
50251_at				AA524283	2.5	2	2.6
74600_g_at	U6 snRNA-associated Sm-like protein LSm8			AI978701	2	1.7	2.6
52932_at	kangai 1 (suppression of tumorigenicity 6, prostate; CD82 antigen (R2 leukocyte antigen, antigen detected by monoclonal and antibody IA4))	KAI1		W51778	3.3	2.9	2.6

Figure 13XXXXXXX

49017_at	hypothetical protein	AA524063	1.8	1.3	2.6
64427_at		AA534724	2.3	0.9	2.6
44694_at	AD-003 protein	AI149096	2.2	1.5	2.6
49024_at		AI078502	2.4	1.5	2.6
53802_at		AA151838	2.6	1.8	2.6
64087_i_at	hypothetical protein FLJ12171	AI022065	2.6	1.9	2.6
46586_at		AI417830	1.4	1.2	2.6
54023_at		AA703247	2.4	2.3	2.6
51937_at		AI376573	2.2	2	2.6
44075_at	p53-induced protein	AA173227	1.4	1.3	2.6
43499_at	tropomyosin 1 (alpha)	N76371	3	2.6	2.6
57836_at	cathepsin D (lysosomal aspartyl protease)	AI541347	1.6	1.9	2.6
55059_at	cytokine inducible SH2-containing protein	AW052069	2.2	0.5	2.6
48005_at		AI862553	2.8	2.8	2.6
41402_at	DKFZP564O0823 protein	AL080121	0	2.5	2.5
41406_at	hypothetical protein FLJ21919	AL080172	2.5	2.1	2.5
39329_at	actinin, alpha 1	X15804	1.5	1.5	2.5
32135_at	sterol regulatory element binding transcription factor 1	U00968	2.4	1.4	2.5
38530_at		W26021	1	1.4	2.5
40905_s_at	DKFZP566J153 protein	AL050369	1.3	0.6	2.5
1388_g_at	vitamin D (1,25-dihydroxyvitamin D3) receptor	J03258	1.9	0.9	2.5
31901_at	potassium voltage-gated channel, shaker-related subfamily, beta member 2	AF044253	2.6	1.6	2.5
34882_at	nucleolar protein (KKE/D repeat)	Y12065	2.6	2.2	2.5
35758_at	RuvB (E coli homolog)-like 2	AB024301	3.4	2	2.5
37740_r_at	solute carrier family 25 (mitochondrial carrier; adenine nucleotide translocator), member 5	J02683	2.5	1.8	2.5
40758_at	immature colon carcinoma transcript 1	X81788	2.5	1.3	2.5
91384_at	rat guanine nucleotide dissociation stimulator	AA147088	2.3	2	2.5
43843_s_at	eukaryotic translation initiation factor 3, subunit 6 (48kD)	AW015717	1.4	1.6	2.5
56458_at		Z99408	2.7	1.4	2.5
46223_at	HSPC025	AI003066	2.4	2.2	2.5
75129_at	myosin IE	AW016142	2.3	1.3	2.5
88767_at		AW008519	2.5	2.5	2.5
64148_at		AW016356	2.6	2.2	2.5

Figure 13YYYYYY

59504_at			AA932206	2.6	2.6	2.5
45317_at			H06725	2.1	1.9	2.5
91825_at			AI375222	2.5	2.5	2.5
48839_s_at			R37261	1.4	1	2.5
48781_s_at			AI992128	3.3	2	2.5
56229_at		hypothetical protein FLJ10349	AI018507	0.6	1.9	2.5
70694_at			AL045971	2.3	2.5	2.5
58774_at		eukaryotic translation initiation factor 4 gamma, 1	AI380979	2.4	1.8	2.5
43528_at			AW003351	2	2.3	2.5
50198_at			AI208272	2.5	2.5	2.5
58784_s_at		GMPR2 for guanosine monophosphate reductase isolog	AI984180	2.1	1.5	2.5
45729_at		KIAA1219 protein	AA156777	2.4	1.8	2.5
45732_at			AL045545	2.1	1.3	2.5
43967_at		ATP-dependant interferon response protein 1	AI826411	3.2	0	2.5
83558_l_at			AI809148	2	1.4	2.5
64495_f_at		SCAN domain-containing 1	AW003586	1.8	2	2.5
51169_at		peptidylprolyl isomerase (cyclophilin)-like 1	AA524353	2.3	2.2	2.5
45828_at		hypothetical protein FLJ10241	AI768100	2.9	1.5	2.5
44043_at			AW004842	2.3	1	2.5
49401_s_at			AI377106	2.3	1.4	2.5
53791_at			AA034010	2.2	1.6	2.5
44605_at			AA521035	2.4	1.7	2.5
45149_at		suppressor of variegation 3-9 (Drosophila) homolog 1	AA604816	2.5	2.2	2.5
52092_at			AA497123	2	1.3	2.5
45603_at		hypothetical protein FLJ20979	AA534993	1.2	1.9	2.5
53660_at			AI240194	2.8	0.7	2.5
46747_s_at		HSPC028 protein	AI460118	1.8	1.9	2.5
46605_at			D20715	1.6	1.4	2.5
74796_at		ralA binding protein 1	T72617	2.6	1.6	2.5
74541_at		low density lipoprotein-related protein 1 (alpha-2-macroglobulin receptor)	AI969879	2.6	1.7	2.5
57960_at			AI949456	1	1.8	2.5
47627_at		cytochrome P540, subfamily IIS, polypeptide 1	AI445492	2.5	2.5	2.5

Figure 13ZZZZZZZ

affyID	Gene Name	Gene Symbol	GBA	Score_Ecoli	Score_Candida	Score_Influenza
909_g_at	interferon-induced protein with tetratricopeptide repeats 2	IFIT2	M14660	320.5	145.5	573.3
908_at	interferon-induced protein with tetratricopeptide repeats 2	IFIT2	M14660	22.9	34	384.9
39945_at	fibroblast activation protein, alpha	FAP	U09278	1.1	1	279.2
1666_at	interferon, type 1, cluster	IFN1@	J00210	2.7	0	191.9
269_at	2'5'oligoadenylate synthetase-like	OASL	L40387	19.8	6.9	119.4
36892_at	integrin, alpha 7	ITGA7	AF032108	0.5	0	92.6
47013_at			AA121732	0.1	3.2	86.3
62493_at			W68034	1.7	0	81.3
47634_at			AW052044	0	0	79.8
44399_at			H11732	1.6	2.1	75.4
44936_at			T93893	3.6	10.2	59
79721_at			AI733197	16.5	11.6	56.6
91973_at			AI470600	0.7	0.1	53.5
39703_at	ubiquitously transcribed tetratricopeptide repeat gene, X chromosome	UTX	AF000993	10.2	4.6	48.9
32065_at	cAMP responsive element modulator	CREM	S68134	7.7	8.5	47.4
60322_at			AI040220	0.8	1.9	47.3
48080_at	hypothetical protein FLJ20037		AI566481	10.4	12.6	45.7
1787_at	cyclin-dependent kinase inhibitor 1C (p57, Kip2)	CDKN1C	U22398	0.3	5.2	45.6
32052_at	hemoglobin, beta	HBB	L48215	0.8	0.7	45.1
75093_at			AI807000	1.5	1	45.1
84717_at			AI307804	0.5	0.7	45.1
34702_f_at	endogenous retroviral protease		M27826	4.3	0	43.9
90048_at	H3 histone, family 3B (H3.3B)	H3F3B	AA609509	0.4	0	42.7
33352_at	H2A histone family, member Q	H2AFQ	X57985	8.1	11.8	37.7
37623_at	nuclear receptor subfamily 4, group A, member 2	NR4A2	X75918	0.7	0.4	37.7
42048_at			AA897644	4.4	2.5	37.2
82094_i_at			AI810266	12.4	6.4	36.3
32182_at	KIAA0965 protein		AB023182	3.6	0	35.9
36085_at	guanine nucleotide binding protein (G protein), alpha transducing activity polypeptide 1	GNAT1	X63749	6	2.4	35.8
74402_at			AI924226	0	0.9	35.7
56608_at	endogenous retroviral protease		AW007800	2.7	1.6	35.5

Figure 14A

40757_at	granzyme A (granzyme 1, cytotoxic T-lymphocyte-associated serine esterase 3)		GZMA	M18737	5.6	1.3	23.9
81908_at				AI650444	0	0.8	23.9
39726_at	glucagon		GCG	J04040	1.1	0	23.6
63607_at	hypothetical protein FLJ20360			AI923841	0	0.7	23.3
59451_at				AI800640	5.3	6.2	23.2
65615_at	KIAA0129 gene product			AA843533	3.2	0.8	22.9
44057_at	hypothetical protein FLJ10633			AA708740	0	0	22.9
67846_at				AI217382	4.8	0.6	22.9
43197_at				AA863064	0.9	0.9	22.8
32587_at	butyrate response factor 2 (EGF-response factor 2)		BRF2	U07802	2	0	22.2
87425_at				AI239840	8.2	0.5	22.2
46926_at				R61533	0	0.5	22
63749_at	KIAA1610 protein			AA411556	2	0	22
63368_at				W81119	0	4.8	22
46829_at				N49940	0.8	3.2	21.4
55940_at				AI701480	0	5.2	21.4
71370_f_at				T55845	0	0	21.2
35849_at	phosphatidylserine receptor			AB011157	0	0.9	21
64683_at				AI681906	1.3	0.1	20.7
47643_at				H10322	0.5	2	20.7
51141_at	protein kinase domains containing protein similar to phosphoprotein C8FW			AI949781	1.3	7.1	20.5
37060_at				U79289	1.5	0.9	20.4
58867_s_at	immunoglobulin lambda locus		IGL@	AI872510	3	2.9	20.2
34595_at	myosin, heavy polypeptide-like (110kD)		MYHL	AF105424	10.9	1.8	20
60812_at	pyruvate dehydrogenase phosphatase			AI625803	0	1	19.9
33273_f_at	immunoglobulin lambda locus		IGL@	X57809	0.5	1.5	19.7
41582_at	advinin			AF041449	1.7	0	19.5
65031_at				AI142111	0.7	1.1	19.1
34460_at	peripheral benzodiazepine receptor-associated protein 1			AB014512	1.9	0.2	19
78599_at				W52073	0.2	2.6	18.9
43373_s_at				AA176915	0	0.1	18.7
56969_f_at				AI658703	8.2	3.1	18.6
48416_at	hypothetical protein HDCMC04P			AA082707	0	0	18.5

Figure 14C

56103_at				H48444	0.3	0.6	18.5
56244_at				H64595	0.6	0.6	18.5
48357_at	ribosomal protein S28		RPS28	AA968694	0.2	5	18.1
91655_f_at				AI419764	3.1	3.5	18.1
43186_at				AA394226	7.3	4.1	18
40199_at	msh (Drosophila) homeo box homolog 1 (formerly homeo box 7)		MSX1	M97676	1.5	0	17.8
86419_f_at				AA837464	2.1	0.3	17.8
59403_at	coatomer protein complex, subunit beta		COPB	AA402414	1	1.6	17.7
50049_f_at				AA045236	0.7	0.6	17.5
35817_at	myelin basic protein		MBP	M13577	2.1	2	17.4
90375_at				N25135	0	0	17.2
52989_at				3143817	0.3	0.1	17.2
58161_at				AA813728	9.7	1.5	17.1
58889_at				R45627	0.2	0	17
40390_at	serine dehydratase		SDS	J05037	3.7	5.2	16.9
40686_at	neuromedin B		NMB	AI985272	2.9	3.2	16.9
60001_at				AA405241	0.6	2.4	16.9
35224_at				AF070569	7.9	1.9	16.8
35868_at	advanced glycosylation end product-specific receptor		AGER	M91211	3.2	1	16.7
34445_at	KIAA0471 gene product			AB007940	1.2	6.4	16.7
46867_at				AI659243	0	0.7	16.2
41277_at	sin3-associated polypeptide, 18kD		SAP18	AW021542	0.7	1	16.1
81061_at				AI978710	0.5	0.9	15.9
36423_at	p8 protein (candidate of metastasis 1)			W47047	3.1	1	15.6
56307_at				AI862477	0	2.3	15.6
61074_at				AI817642	0.4	0	15.4
36760_at				U79277	1.2	0.9	15.3
65008_g_at	splicing factor (CC1.3)			AI539492	2.1	1.1	15.3
60190_at				AI668659	1.8	1.7	15.3
51032_at	proteasome (prosome, macropain) 26S subunit, ATPase, 2		PSMC2	AI754675	0	0	15.3
36097_at	immediate early protein			M62831	6.2	1.6	15.2
32344_r_at	potassium voltage-gated channel, shaker-related subfamily, member 4		KCNA4	M55514	0.3	0.8	15
44594_at	CGI-112 protein			AA570477	1	2.2	15

Figure 14D

40434_at	podocalyxin-like	PODXL	U97519	3.7	1.4	14.9
76540_at			AA350911	0.8	0	14.8
48259_at			AI123757	3.7	2.4	14.7
49258_at	hypothetical protein DKFZp761B2423		AW007698	10	1	14.7
636_at			L43338	1.1	0.2	14.7
54579_at			AI799534	0.1	5.9	14.6
74681_at			AI018220	3.6	1.4	14.5
48978_at			AI769281	1.4	1.8	14.5
38265_at	hypothetical protein DKFZp761B2423		AI538172	7.7	0.8	14.4
39364_s_at	protein phosphatase 1, regulatory (inhibitor) subunit 5	PPP1R5	Y18207	1.7	0.3	14.1
36059_at	low density lipoprotein receptor-related protein 4	LRP4	AB011540	1	1	14
55712_at			AI821417	0.4	1.7	14
35488_at	small nuclear RNA activating complex, polypeptide 1, 43kD	SNAPC1	U44754	3.4	0.8	13.9
823_at	small inducible cytokine subfamily D (Cys-X3-Cys), member 1 (fractalkine, neurotactin)	SCYD1	U84487	3.9	1.1	13.9
1089_i_at			M64936	11.1	3.3	13.9
47032_at			AI885390	1.5	4.3	13.8
37249_at	phosphodiesterase 8B	PDE8B	AF079529	1.1	0	13.5
38017_at	CD79A antigen (immunoglobulin-associated alpha)	CD79A	U05259	0.3	0.5	13.5
143_s_at	TATA box binding protein (TBP)-associated factor, RNA polymerase II, D, 100kD	TAF2D	U75309	1.3	0	13.5
58182_r_at			AA814034	0	1.8	13.5
41052_s_at	calcium channel, voltage-dependent, P/Q type, alpha 1A subunit	CACNA1A	U79666	0.3	3	13.4
84064_at			AI365233	0.8	1.3	13.4
64194_at	hypothetical protein FLJ10903		H14920	0	0.2	13.4
63746_i_at			AA279977	2.3	0	13.4
89589_f_at			R05527	0	0.1	13.2
36214_at	Kruppel-like factor 4 (gut)	KLF4	U70663	0.2	0	13.1
88861_at			AI680122	0.7	0.9	13.1
62449_i_at	hypothetical protein DKFZp761C169		R82825	2.4	2.7	13.1
45178_at	RAP1A, member of RAS oncogene family	RAP1A	AI476340	0	0.1	13
40479_at	FYN oncogene related to SRC, FGR, YES	FYN	Z97989	0.3	0	12.7
38156_at	MyoD family inhibitor	MDFI	U78313	0.4	0	12.6
31889_at	melan-A	MLANA	U06452	4.2	0.3	12.6

Figure 14E

72959_at				T93073	1.4	3.2	12.5
1894_f_at				L27065	0.5	0.4	12.5
51388_at				AA827683	0	0	12.4
1435_f_at			IFNA21	V00540	0	0.5	12.3
58550_at			interferon, alpha 21	AA081874	0.9	0.2	12.3
43986_at			similar to aspartate beta hydroxylase (ASPH)	H17272	0.3	0.5	12.3
53567_at				AW003119	0	0.2	12.3
54648_at				AA810864	0.2	1.6	12.2
37112_at			chromosome 6 open reading frame 32	AB002384	0.2	0	11.9
41665_at			PCF11p homolog	AB020631	0	0.1	11.9
39636_at				AL079294	0.7	0.8	11.9
44241_r_at				T92908	0	1.6	11.8
57942_at				AA282811	1.5	0.6	11.8
37228_at			polo (Drosophila)-like kinase	U01038	0.3	0	11.7
64566_at				N54523	1.1	2	11.7
42770_at				T86276	1.1	1.5	11.6
56141_at				AI566128	2.3	0.4	11.6
81057_at				AA085091	4.2	1.3	11.5
51022_at				AI377043	0	2.3	11.5
54243_at				AA780058	1.5	1.1	11.5
37909_at			laminin, alpha 3 (nicein (150kD), kalinin (165kD), BM600 (150kD), epilegrin)	L34155	0.6	1.2	11.4
71530_at				AI936520	0.2	1.3	11.4
47120_at				AW005696	0	0	11.4
39953_i_at			KIAA0628 gene product	AB014528	1.2	0.1	11.3
40667_at			CD6 antigen	X60992	1.5	2.3	11.3
60514_at			LATS (large tumor suppressor, Drosophila) homolog 2	U69566	3.3	0.4	11.3
34286_at			breast carcinoma amplified sequence 2	AB020623	0.7	0.8	11.2
32415_at			interferon, alpha 5	V00541	0.4	2.4	11.2
35955_at			cytochrome c-like antigen	S80864	0.7	0.2	11.1
40091_at			B-cell CLL/lymphoma 6 (zinc finger protein 51)	U00115	0	0.9	11
50652_at				N53764	0.1	0.3	11
55631_at			KIAA1414 protein	AI499339	0.3	0	11
79089_at				AI797610	0	1.5	10.9
49788_at				AA102395	0.3	1.9	10.9

Figure 14F

39846_at	cathepsin F	CTSF	AF071748	0.1	1.7	10.8
41854_at	v-myb avian myeloblastosis viral oncogene homolog	MYB	M13666	0.8	0	10.8
33118_at	interferon-related developmental regulator 2	IFRD2	U73167	0	0.6	10.8
56213_i_at			AA046067	1.4	1.4	10.8
85344_f_at			AW024994	0	0	10.8
1575_at	ATP-binding cassette, sub-family B (MDR/TAP), member 1	ABCB1	M14758	0	0.9	10.7
38041_at	UDP-N-acetyl-alpha-D-galactosamine:polypeptide N-acetylgalactosaminyltransferase 1 (GalNAc-T1)	GALNT1	U41514	0	0	10.7
57642_at			AA005336	1.6	0.9	10.7
59220_at			AA922861	0.6	1.6	10.7
48672_at			A1817153	1.2	0.2	10.7
52985_at			3075759	0.1	0.6	10.7
90878_at			T79248	1.1	1.9	10.6
1541_f_at	interferon, alpha 6	IFNA6	X02958	0.4	0	10.4
36315_i_at	KIAA0919 protein		AB023136	0.5	0.4	10.4
61150_at			AA894558	0.5	0.3	10.4
65331_at			AW009884	1.9	2.5	10.4
71791_at			A1347488	0	0.3	10.4
32882_at	sterol O-acyltransferase 2	SOAT2	AF059203	1.2	0	10.3
41784_at	DKFZP564B0769 protein		AL080186	0	0.3	10.2
602_s_at	hydroxysteroid (17-beta) dehydrogenase 1	HSD17B1	M29037	0	0.7	10.2
66524_at			AA830101	0.4	0.5	10.2
44947_at			AA047365	1.7	2.3	10.2
53595_at	AD-015 protein		N39009	0.6	3.4	10.2
40457_at	splicing factor, arginine/serine-rich 3	SFRS3	AF038250	0.4	0.7	10.1
38701_at	ubiquitin-conjugating enzyme E2L 3	UBE2L3	AJ000519	0	2.4	10
32583_at	v-jun avian sarcoma virus 17 oncogene homolog	JUN	J04111	2.1	2.7	9.9
37896_at	trefoil factor 3 (intestinal)	TFF3	A474125	2.3	0.1	9.9
33012_at	tumor necrosis factor (ligand) superfamily, member 8	TNFSF8	L09753	0.5	0	9.8
36343_at	toll-like 2	TLL2	AB023149	2.5	0.7	9.8
71984_at			W04199	0	0.8	9.8
49372_at			N67059	0.2	0.1	9.7
45427_at			R51079	0.4	0.9	9.7
43760_at			AA397579	1.4	0.6	9.7
59479_at			W44692	1.2	0.8	9.6

Figure 14G

58459_s_at			AA150805	0	0.9	9.6
58457_f_at			A1819225	0.2	0	9.6
36866_at	testis-specific ankyrin motif containing protein		D78334	2.1	2.4	9.5
54501_at			A1253196	0	0	9.5
39650_s_at	KIAA0435 gene product		AB007895	1.6	1.9	9.4
36053_at	cyclin-dependent kinase inhibitor 2C (p18, inhibits CDK4)	CDKN2C	AF041248	0	0	9.4
44122_at	transcriptional regulator protein		A1074024	2.4	2.7	9.4
44212_at			AA167191	0.4	0	9.4
1222_at	PCTAIRE protein kinase 2	PCTK2	X66360	0.4	0.1	9.3
796_i_at	cyclin-dependent kinase-like 1 (CDC2-related kinase)	CDKL1	X66358	0	0	9.3
1951_at	angiopoietin 2	ANGPT2	AF004327	0.5	0	9.2
31924_at	soluble adenylyl cyclase		AL035122	1.7	0	9.2
32648_at	delta-like homolog (Drosophila)	DLK1	U15979	0.6	0	9.2
57658_at			U55989	1	0	9.2
51609_r_at			AW051767	0	1	9.2
34594_at	related to the N terminus of tre		D13644	0.2	0.8	9.1
63968_at			A1979124	1.3	0	9.1
42056_at			AA282541	1.8	1.6	9.1
41455_at	KIAA0712 gene product		AB018255	0.7	0.1	9
32319_at	tumor necrosis factor (ligand) superfamily, member 4 (tax- transcriptionally activated glycoprotein 1, 34kD)	TNFSF4	AL022310	1.2	1.4	9
1452_at	LIM domain only 4	LMO4	U24576	5.7	1.5	9
33952_at	zinc finger protein zfp47		U71601	0.4	0.8	9
37578_at	actin filament associated protein		D25248	0.7	0.7	9
51307_at			A1589165	0.2	1.9	9
48417_r_at	hypothetical protein PRO2849		AA636101	0	0.2	9
47823_r_at			A1452659	0	0	9
61789_i_at	proteasome (prosome, macropain) 26S subunit, non-ATPase, 11	PSMD11	H05559	0.3	0.2	9
33365_at	KIAA0945 protein		AB023162	0.2	0.2	8.9
59870_at			A1003552	2.3	1.1	8.9
91329_at			A1669284	0	0	8.9
80897_at			R63836	1.8	1.7	8.9
34057_at	solute carrier family 28 (sodium-coupled nucleoside transporter), member 2	SLC28A2	U84392	2	0	8.8

Figure 14H

1048_at	retinoid X receptor, gamma	RXRG	U38480	1.7	0.8	8.8
56459_at			AI821762	0.9	0.6	8.8
52425_at	KIAA1376 protein		AA164405	1.4	0	8.8
84483_f_at			AI459092	1.7	0.9	8.8
32254_at	vesicle-associated membrane protein 2 (synaptobrevin 2)	VAMP2	AL050223	1.8	1.7	8.7
45889_at			AW024527	0	0	8.7
40552_s_at			AL049987	0	0	8.6
59414_at			AI480103	0.8	0	8.6
43958_at			AI499157	0	0.9	8.6
43031_at			AL045916	0	1.8	8.6
1241_at	protein tyrosine phosphatase type IVA, member 2	PTP4A2	U14603	0	0.5	8.5
79512_at			AI248270	3	2.5	8.5
55974_at			AI732621	0.8	0.9	8.5
40641_at	TBP-associated factor 172		AF038362	0	1.9	8.4
33455_at	aldolase B, fructose-bisphosphate	ALDOB	X02747	0	1.1	8.4
40738_at	CD2 antigen (p50), sheep red blood cell receptor	CD2	M16336	0.8	1	8.4
60723_at			AA744550	1	0.9	8.4
84319_at			AI684171	0	0	8.4
33489_at	trefoil factor 2 (spasmolytic protein 1)	TFF2	U47292	1	0	8.4
44602_at			AA156238	0.3	1	8.4
52750_at			AI632711	1.6	0.4	8.4
31970_r_at	protein phosphatase, EF hand calcium-binding domain 2	PPEF2	AF023456	0.7	0.6	8.3
59347_at			R25153	0	1.5	8.3
67975_at			AA934126	0.8	0.9	8.3
85630_at			AI086384	0	0.3	8.3
42764_at			AI459108	0.1	0.1	8.2
58370_at	hypothetical protein DKFZp761A052		AW003291	0.2	0.4	8.2
53468_at			AI149508	0.2	0.3	8.2
64394_at	cadherin 12, type 2 (N-cadherin 2)	CDH12	AI476284	0.4	0.6	8.2
33538_at	myelin gene expression factor 2		H96671	1.1	0.5	8.1
1826_at	ras homolog gene family, member B	ARHB	M12174	0	0	8
32288_r_at	killer cell lectin-like receptor subfamily C, member 3	KLRC3	AJ001685	0.6	0.5	8
34573_at	ephrin-A3	EFNA3	U14187	0.6	0	8
72361_at			AI885066	0	0	8
80940_f_at			AI147061	0.1	0.2	8

Figure 14I

629_at	interleukin 9 receptor	IL9R	L39064	0.2	0.2	7.9
89024_at			A1720099	0.9	0.3	7.9
34502_g_at	runt-related transcription factor 2	RUNX2	L40992	0	0	7.8
39691_at	SH3-containing protein SH3GLB1		AB007960	0	0.7	7.8
33295_at	Duffy blood group	FY	X85785	0.5	0.3	7.8
37551_at	KIAA0211 gene product		D86966	0.5	0.8	7.8
83013_i_at			A1400387	1.1	0.8	7.8
69708_at			A1689617	1.6	1.6	7.8
41190_at	tumor necrosis factor receptor superfamily, member 12 (translocating chain-association membrane protein)	TNFRSF12	U83598	0.7	0	7.7
62857_at			A1784540	1.3	0.4	7.7
71584_at	X-box binding protein 1	XBPI	AA503434	2.5	0	7.7
82126_at			AA831700	0.8	0.1	7.7
64460_at	M-phase phosphoprotein homolog		A1823360	0.9	1	7.7
47018_at			A1168404	0	0.8	7.7
36050_at	zinc finger protein 313	ZNF313	AL031685	0.6	0.5	7.6
40681_at	integrin, beta-like 1 (with EGF-like repeat domains)	ITGBL1	AB008375	1	0.4	7.6
53862_at			AA447888	0.1	0.7	7.6
50853_at			A1299719	0.8	0.4	7.6
53576_at			AA041298	0.2	0.9	7.6
1317_at	macrophage stimulating 1 receptor (c-met-related tyrosine kinase)	MST1R	X70040	0.8	2	7.5
38772_at	cysteine-rich, angiogenic inducer, 61	CYR61	Y11307	0.4	1	7.5
38319_at	CD3D antigen, delta polypeptide (TiT3 complex)	CD3D	AA919102	0.6	0.4	7.5
66366_at			A1695056	0	0.6	7.5
82251_at			A1819034	0.5	0.7	7.5
37161_at			W28948	0.7	0.4	7.5
54125_at			30383358	0.3	0.5	7.5
72546_at			A1033943	1.3	1.4	7.5
38187_at	N-acetyltransferase 1 (arylamine N-acetyltransferase)	NAT1	D90041	0.5	0.5	7.4
37681_i_at	matrin 3	MATR3	AB018266	1.4	0.3	7.4
41107_at	syntaphilin		AB002372	0.8	0	7.4
67068_at			A1298437	0.3	2.4	7.4
71483_at			A1937154	0.1	0.8	7.4
37996_s_at	dystrophin myotonic-protein kinase	DMPK	L08835	0	0	7.3

Figure 14J

2021_s_at	cyclin E1	CCNE1	M73812	0.4	0	7.3
45710_at	DNA segment, single copy, probe pH4 (transforming sequence, thyroid-1,	D10S170	H15313	0	0	7.3
92101_s_at	ubiquitin A-52 residue ribosomal protein fusion product 1	UBA52	AI802408	0	0.6	7.3
34931_at	KIAA0940 protein		AB023157	0.8	0.9	7.2
40428_i_at			AW043812	0	0.3	7.2
82383_r_at			AI215733	0.5	0	7.2
45220_at			AA150501	0.6	2.2	7.2
1280_i_at			Z25424	0.1	0	7.2
1937_at			M26460	0.8	0	7.2
1252_at	DNA segment, single copy probe LNS-CAI/LNS-CAII (deleted in polyposis		M73547	0	0.7	7.1
37925_r_at	apolipoprotein M		AJ245434	0	0.8	7.1
33922_at	PR domain containing 2, with ZNF domain	PRDM2	U17838	0	0	7.1
844_at	protein phosphatase 1, regulatory (inhibitor) subunit 1A	PPP1R1A	U48707	0.6	0.5	7.1
39476_at			Z49995	0.9	0.6	7.1
38127_at	syndecan 1	SDC1	Z48199	0.1	0.1	7
62624_at			AL043110	0	1	7
78727_at			AA425815	0	0	7
37130_g_at	neuropeptide FF-amide peptide precursor	NPFF	AI936758	0.4	0.4	6.9
34175_r_at			AL049242	0.6	0	6.9
32793_at	T cell receptor beta locus	TRB@	X00437	0.7	0.3	6.9
35359_at	KIAA0235 protein		D87078	0.8	0.1	6.9
67857_at			AA425402	0.2	0.3	6.9
57101_at			AI242202	0.3	1.2	6.9
80665_r_at			AI963771	2.1	1.9	6.9
56683_i_at			AA639553	1.6	2.3	6.9
61180_f_at			AA024934	0.7	0	6.9
176_at	protein phosphatase 2, regulatory subunit B (B56), gamma isoform	PPP2R5C	U37352	0	0	6.8
33881_at	fatty acid:Coenzyme A ligase, long chain 3	FACL3	AA977580	0.3	0	6.8
35568_at	adenylate cyclase activating polypeptide 1 (pituitary) receptor type I	ADCYAP1R1	D17516	1.7	2.1	6.8
36949_at	casein kinase 1, delta	CSNK1D	U29171	0.6	0.7	6.8
38284_at	KIAA0304 gene product		AJ007041	2.1	1.7	6.8

Figure 14K

50545_at			AA947458	1.7	1.9	6.8
78786_at			AI091533	1.6	2.7	6.8
59357_at			AI433468	0	0	6.8
36757_at	H2A histone family, member C	H2AFC	AL009179	0.5	0.4	6.7
91032_at			AI791640	0.3	0	6.7
80124_at			AA643238	0.6	0	6.7
86332_at			AI523641	1	0.1	6.7
43479_at			AA082768	0.5	0.2	6.7
61321_f_at			AA534354	0	0.8	6.7
31508_at	upregulated by 1,25-dihydroxyvitamin D-3		S73591	0	0.1	6.6
1044_s_at	E2F transcription factor 5, p130-binding	E2F5	U31556	2.3	0.4	6.6
33872_at	latrophilin		AB018329	1.5	0	6.6
793_at	placental growth factor, vascular endothelial growth factor-related protein	PGF	X54936	0.6	2.9	6.6
1217_g_at	protein kinase C, beta 1	PRKCB1	X07109	1	0.5	6.6
67262_at			AI378639	0.5	0	6.6
50352_at	hypothetical protein FLJ10652		AA151706	0	0.7	6.6
64672_at			AI339915	0	0.6	6.6
46454_at			AA401446	0.1	0.4	6.6
51938_at			N33202	0.4	0	6.6
31559_at	solute carrier family 13 (sodium-dependent dicarboxylate transporter), member 2	SLC13A2	U26209	0.8	0	6.5
34237_at	HBS1 (S. cerevisiae)-like	HBS1L	AB028961	0.9	0.6	6.5
34708_at	ficollin (collagen/fibrinogen domain-containing) 3 (Hakata antigen)	FCN3	D88587	0.7	0.6	6.5
56930_at	integrin beta 1 binding protein (melusin) 2	ITGB1BP2	AI568769	1.1	1.1	6.5
55540_at			AI806973	2.8	1.2	6.5
55542_at			AI735416	0	0.2	6.5
77657_f_at			AI371706	0.2	0.4	6.5
1373_at	transcription factor 3 (E2A immunoglobulin enhancer binding factors E12/E47)	TCF3	M31523	0	0.3	6.4
32681_at	solute carrier family 9 (sodium/hydrogen exchanger), isoform 1 (antiporter, Na+/H+, amiloride sensitive)	SLC9A1	S68616	0	0.9	6.4
40576_f_at	heterogeneous nuclear ribonucleoprotein D-like	HNRNPD	D89678	0	0	6.4
40041_at	highly expressed in cancer, rich in leucine heptad repeats		AF017790	0	0.9	6.4

Figure 14L

48078_at			W04245	0	0	6.4
88916_at			AI703450	0	0	6.4
74508_at			AW015038	0.4	1.4	6.4
44667_at			AI422758	0	0	6.4
72496_f_at			AA552962	1	0.4	6.4
33399_at	ribosomal protein S6	RPS6	AA142942	0	1.1	6.3
39609_at	single-minded (Drosophila) homolog 2	SIM2	U80457	0.4	0.2	6.3
39940_at			AL080094	0.9	0	6.3
47443_at			AI268054	0.5	0.8	6.3
70085_f_at			AI817150	1	0.3	6.3
49633_at	hypothetical protein HES6		N21131	0	0.5	6.3
70198_f_at			AI821980	1.2	0.3	6.3
82253_i_at			AA663583	1.3	0	6.3
86760_f_at			AI096457	0	0.5	6.3
32106_at	serine (or cysteine) proteinase inhibitor, clade A (alpha-1 antiproteinase, antitrypsin), member 4	SERPINA4	L28101	0.5	0.5	6.2
40539_at	myosin IXB	MYO9B	U42391	1	0.7	6.2
41196_at	karyopherin (importin) beta 1	KPNB1	L38951	0	0	6.2
33175_at			AA156237	0.9	0	6.2
35829_at	immunoglobulin superfamily, member 4	IGSF4	AL080181	0.5	0	6.2
51716_r_at	complement component 5	C5	T71645	0.2	0.1	6.2
50062_at			AA622412	0.6	0.2	6.2
62008_at			AI935871	0.8	0.2	6.2
48961_at			N58188	0.9	0.5	6.2
52309_at			2969294	0.7	0.2	6.2
1552_i_at	cytochrome P450, subfamily IIA (phenobarbital-inducible), polypeptide 13	CYP2A13	U22028	1.8	0.8	6.1
39563_at	KIAA0268 protein		D87742	0.4	0.2	6.1
34621_at	keratin 2A (epidermal ichthyosis bullosa of Siemens)	KRT2A	AF019084	0.6	0.1	6.1
55010_g_at			AA451732	0.5	0.2	6.1
81965_at			H19232	1	0.9	6.1
61856_at			AI634650	0.2	0	6.1
84131_at			AI808593	0	0.9	6.1
56939_at			AA399009	2.9	2.6	6.1
86817_at			AF150371	1.2	1	6.1

Figure 14M

63458_r_at				AA022988	0	1	6.1
38747_at	CD34 antigen	CD34		M81945	0.7	0.2	6.1
92138_at				A1345255	0.3	0	6.1
79933_at				N33374	0.9	2.1	6.1
1974_s_at	tumor protein p53 (Li-Fraumeni syndrome)	TP53		X02469	0.7	0	6
38901_at	ubiquitin specific protease 19	USP19		AB020698	0	0.3	6
39951_at	plastin 1 (l isoform)	PLS1		L20826	1.9	0.7	6
88770_r_at				A1694231	0.6	0	6
42757_at				AA424339	0	0.6	6
48435_at				A1796988	0	0.8	6
65108_at				A1346432	0.2	1.5	6
51381_at				A1480091	0	0	6
40487_at				W26634	0	0	5.9
768_at	karyopherin alpha 5 (importin alpha 6)	KPNA5		AF005361	1.2	0	5.9
34231_at	histone acetyltransferase			AF074606	0	0.7	5.9
37576_at	Purkinje cell protein 4	PCP4		U52969	0.3	0.4	5.9
47059_at	splicing factor, arginine/serine-rich 11	SFRS11		AI002238	0	0	5.9
45522_at				R41958	0	0.4	5.9
88177_f_at				AI492518	0.8	0.4	5.9
57714_at				AA429113	0	0.8	5.9
990_at	fms-related tyrosine kinase 1 (vascular endothelial growth factor/vascular permeability factor receptor)	FLT1		X51602	0.2	0.4	5.8
38521_at	CD22 antigen	CD22		X59350	0.4	0.6	5.8
38561_at	down-regulated in metastasis			N58318	1.2	0.4	5.8
36061_at				AF009314	0	0.5	5.8
52686_at				AI350227	0.7	0.9	5.8
86831_at				T91952	0.2	0.1	5.8
86865_at				AA009833	0	0	5.8
69774_at				AI024369	1.5	1.7	5.8
1411_at				D16154	0.6	0.4	5.8
524_at	postmeiotic segregation increased (S. cerevisiae) 1	PMS1		U13695	0.9	0.7	5.7
39790_at	ATPase, Ca++ transporting, cardiac muscle, slow twitch 2	ATP2A2		M23115	0.1	0.5	5.7
33359_at	KIAA0768 protein			AB018311	0.8	0.1	5.7
32405_at	KIAA0707 protein			AB014607	1	0	5.7
33515_at	t-complex 10 (a murine tcp homolog)	TCP10		U03399	0	0	5.7

Figure 14N

39199_at				W28661	0.1	0	5.7
752_s_at	DnaJ (Hsp40) homolog, subfamily B, member 1	DNAJB1		D85429	0	0.1	5.7
41018_at	DKFZP564O243 protein			AL050015	0.7	0.8	5.7
42301_at				AA444100	0.3	0.6	5.7
66550_at				AA972106	0.2	0	5.7
50879_at				AI675751	0.1	1.6	5.7
46944_at				AI276680	0.6	1	5.7
64221_at				H27234	0	0	5.7
62391_f_at				AA592983	0	0.3	5.7
33010_at	solute carrier family 5 (neutral amino acid transporters, system A), member 4	SLC5A4		AL008723	0	0.4	5.6
1361_at	telomeric repeat binding factor (NIMA-interacting) 1	TERF1		U40705	0	1	5.6
33080_s_at	RAP1, GTPase activating protein 1	RAP1GA1		AB007943	1	0.7	5.6
1022_f_at	interferon, alpha 14	IFNA14		V00542	0.1	0	5.6
86405_at				Z38166	0	0.4	5.6
34572_at	carbonic anhydrase VII	CA7		M76424	1	0.9	5.6
46391_i_at				AI761712	0.2	1	5.6
46103_at	MEK partner 1			AI335267	0.2	0.6	5.6
32460_at	gamma-aminobutyric acid (GABA) A receptor, beta 2	GABRB2		S67368	0	0	5.5
31586_f_at	immunoglobulin kappa constant	IGKC		X72475	0	0.9	5.5
40582_at	protein tyrosine phosphatase, receptor type, J	PTPRJ		AI806482	0.4	0	5.5
33284_at	myeloperoxidase	MPO		M19507	1.3	0.8	5.5
40291_r_at	sialyltransferase 4A (beta-galactosidase alpha-2,3-sialyltransferase)	SIAT4A		L13972	1.4	0.9	5.5
36813_at	thyroid hormone receptor interactor 13	TRIP13		U96131	0	0.3	5.5
33136_at	ubiquitin-conjugating enzyme E2I (homologous to yeast UBC9)	UBE2I		AL031714	0.1	0.7	5.5
91081_at				AI131053	0.9	0.6	5.5
87820_at				AA287372	1.1	1.5	5.5
67369_at				AA828830	0.1	0.4	5.5
44762_at	mutS (E. coli) homolog 6	MSH6		AI926554	0.6	0.3	5.5
47375_at				AA019836	0.6	0.8	5.5
65951_at	hypothetical protein FLJ20080			AA205862	0.3	0.2	5.5
51113_at	UDP-N-acetyl-alpha-D-galactosamine:polypeptide N-acetylglactosaminyltransferase 1 (GalNAc-T1)	GALNT1		AW021000	0.2	0.5	5.5
50104_at	v-ets avian erythroblastosis virus E26 oncogene homolog 2	ETS2		AI870399	0	0	5.5

Figure 140

62328_at				T92565	0	0	0	5.5
37415_at	ATPase, Class V, type 10B	ATP10B		AB018258	0.3	0	0	5.4
39392_at	glycerophosphate O-acyltransferase	GNPAT		AJ002190	0.5	0	0	5.4
36925_at	heat shock 70kD protein 2	HSPA2		L26336	0	0.2	0	5.4
1231_at	ubiquitin-conjugating enzyme E2B (RAD6 homolog)	UBE2B		M74525	0.8	1	0	5.4
38300_at	frizzled (Drosophila) homolog 1	FZD1		AB017363	0.1	0	0	5.4
65510_f_at				AA034458	0.4	0.4	0	5.4
43177_at				AI732470	0.3	0.1	0	5.4
91067_i_at				AI681736	0.5	0.2	0	5.4
35109_at	KIAA0756 protein			AB018299	0	0.4	0	5.3
33896_at	E1A binding protein p300	EP300		U01877	0.9	0	0	5.3
49292_s_at	ras responsive element binding protein 1	RREB1		AI097330	0	0.1	0	5.3
86470_at				AA884313	0.1	0	0	5.3
88607_at				AI660692	0.9	0.2	0	5.3
45159_r_at				H71807	0.3	0.4	0	5.3
48313_at				AI565705	0	0	0	5.3
91678_at	CGI-14 protein			AI814761	3.3	2.6	0	5.3
55474_at				R32893	0.2	0.4	0	5.3
31668_f_at				W28193	0.8	0.1	0	5.3
86188_f_at				AI733033	0.6	0.6	0	5.3
35066_g_at	fetal hypothetical protein			U84971	0.7	1.5	0	5.2
41692_at	synaptotagmin 1	SYNJ1		AB020717	0	0	0	5.2
39428_at	lymphocyte adaptor protein			AF055581	0.4	0.3	0	5.2
31808_at	inhibitor of growth family, member 3	ING3		AC004537	1.1	0.2	0	5.2
33373_at				AL049951	0.6	0.8	0	5.2
33817_at	heterogeneous nuclear protein similar to rat helix destabilizing protein			S63912	0	0.3	0	5.2
35719_at	KIAA0606 protein; SCN Circadian Oscillatory Protein (SCOP)			AB011178	1	0	0	5.2
58010_at				R27299	0	0.8	0	5.2
89821_at				AI656531	0.7	0	0	5.2
58115_at				AA806110	0.6	0.5	0	5.2
77328_at				AI223339	0	0.7	0	5.2
80861_at				AA424811	0.7	0	0	5.2
68187_at				AA632138	0	0	0	5.2
46929_at				N70317	0	0	0	5.2

Figure 14P

50930_at			AI796010	0	0	5.2
54969_at			AI829724	0	0.9	5.2
42193_r_at			AA434246	0.4	0.2	5.2
34467_g_at	5-hydroxytryptamine (serotonin) receptor 4	HTR4	Y12505	0.6	0.4	5.1
40187_at			AW016815	0.7	1.4	5.1
40067_at	E74-like factor 1 (ets domain transcription factor)	ELF1	M82882	0.5	0.5	5.1
35459_at	regulator of G-protein signalling 13	RGS13	AF030107	0.5	0	5.1
32010_at	hypothetical protein		Z82180	0	0.9	5.1
282_at	M-phase phosphoprotein 1	MPHOSPH1	L16782	0.7	0	5.1
32271_at	FOS-like antigen-1	FOSL1	X16707	0	0	5.1
33842_at	Fanconi anemia, complementation group G	FANCG	AC004472	0	0	5.1
37976_at	Ig superfamily protein		AL034397	0.7	0.5	5.1
41028_at	ryanodine receptor 3	RYR3	AJ001515	0.5	0.1	5.1
42496_at			AA608964	0.7	0.6	5.1
69778_at			AI972712	0	0.7	5.1
51054_r_at	hypothetical protein FLJ10737		W87554	0.1	0.2	5.1
44096_at	desmin	DES	AA527080	0.6	0	5.1
52516_at			R44595	0.2	0.9	5.1
64914_at			AI632224	0.7	0	5.1
52813_at	calumenin	CALU	AA234047	0	0	5.1
31402_at	minor histocompatibility antigen HB-1		AF103884	1	0.4	5
33716_at			N95443	0	0	5
31935_s_at	DEAD/H (Asp-Glu-Ala-Asp/His) box polypeptide 11 (S.cerevisiae CHL1-like helicase)	DDX11	U75968	0.8	0	5
41792_at	ATP-binding cassette, sub-family C (CFTR/MRP), member 8	ABCC8	L78207	0.9	0.6	5
49409_i_at			AA910006	0	0	5
75954_at			AI978606	0	0	5
60924_i_at			H63111	0	0	5
65938_at	MADS box transcription enhancer factor 2, polypeptide C (myocyte enhancer factor 2C)	MEF2C	AL118637	0	0	5
48329_at			AA876002	0.7	0.7	5
53017_at			3144516	0	0	5
74926_r_at			AI252805	1	0.5	5
37140_s_at	ectodermal dysplasia 1, anhidrotic	ED1	AF061193	0.5	0.7	4.9
33011_at	neurotensin receptor 2	NTSR2	Y10148	0.6	0	4.9

Figure 14Q

31890_s_at	zinc finger protein 143 (clone pHZ.1)	ZNF143	AF071771	0.6	0	4.9
1584_at	cell division cycle 25C	CDC25C	M34065	3.1	1.8	4.9
37746_r_at	suppression of tumorigenicity 5	ST5	U15131	0.8	0.1	4.9
418_at	antigen identified by monoclonal antibody Ki-67	MKI67	X65550	0.4	0	4.9
320_at	peroxisomal biogenesis factor 6	PEX6	D83703	0	0.6	4.9
90922_at			AI608696	0.5	0.2	4.9
50475_at			AI679625	0	0	4.9
56258_f_at			AI370094	1	0.2	4.9
92163_at			AA504454	0.8	0	4.9
69668_at			AI744371	0.3	0.8	4.9
65405_at			H67680	0	1	4.9
62740_at			T72588	0.5	0.4	4.9
40445_at	E74-like factor 3 (ets domain transcription factor, epithelial-specific)	ELF3	AF017307	0.8	0	4.8
33387_at	growth arrest-specific 7	GAS7	AB007854	0	0	4.8
37334_at	heterogeneous nuclear ribonucleoprotein A0	HNRPA0	U23803	0	0.1	4.8
197_at	non-metastatic cells 3, protein expressed in	NME3	U29656	0.8	0	4.8
50795_at			AI693592	0.5	0	4.8
49415_at			AI027972	0.7	0.8	4.8
65268_at			AA526937	0.4	0.6	4.8
72264_at			AI948563	0	0.6	4.8
43103_at			AA394147	0.5	0.9	4.8
53274_r_at			R66718	0.8	0.9	4.8
54198_at			AA262451	0.4	0.7	4.8
70656_at			AI827487	0.5	0.8	4.8
73467_f_at			AI799544	0.6	0.7	4.8
38624_at	solute carrier family 12 (potassium/chloride transporters), member 4	SLC12A4	AF054506	1	0	4.7
34940_at			AL080095	1	0	4.7
31408_at	retinal pigment epithelium-derived rhodopsin homolog	RRH	AF012270	0.4	0	4.7
32297_s_at	killer cell lectin-like receptor subfamily C, member 2	KLRC2	AJ001684	0.2	0	4.7
40984_at	gamma tubulin ring complex protein (76p gene)		W28255	0.9	1	4.7
33463_at	xanthene dehydrogenase	XDH	U39487	1	0.6	4.7
41002_at	solute carrier family 16 (monocarboxylic acid transporters), member 5	SLC16A5	U59299	0.6	0.3	4.7

Figure 14R

46462_at			AI822047	0.6	0.4	4.7
58008_at			AA521240	0.4	0.5	4.7
66352_at			AI003755	0.2	0.7	4.7
58366_at			R71447	0	0	4.7
69658_at			AI754641	0.6	0.5	4.7
56681_at			AA628511	0	0.6	4.7
43998_at			AI954729	0	0.6	4.7
65265_r_at			T90778	0	0.7	4.7
59474_at			AI924028	0.5	1	4.7
39866_at		USP22	AB028986	0.6	0.8	4.6
39696_at			AB028974	0.8	0	4.6
40149_at			AL049924	0	0	4.6
32868_at		TGM3	L10386	0.9	0.7	4.6
40631_at		TOB1	D38305	0	0.3	4.6
34779_at		NeuAc-alpha-2,3-Gal-beta-1,3-GalNAc-alpha-2, 6-sialyltransferase alpha2,6-sialyltransferase	R90942	1	0.1	4.6
38442_at		microfibrillar-associated protein 2	U19718	0.4	0	4.6
46790_r_at			AA421236	0.4	0.4	4.6
73870_at			AI700737	0.5	0.6	4.6
35593_at		amine oxidase, copper containing 2 (retina-specific)	AB012943	0.4	0.9	4.6
70246_at			AA630253	1	0.2	4.6
60606_r_at			N68442	0.6	0.5	4.6
45119_at			AI638568	0	0	4.6
60129_f_at			R92776	0.1	0.9	4.6
45590_at			AI673025	0.7	1	4.6
46308_at		a disintegrin-like and metalloprotease (reprolysin type) with thrombospondin type 1 motif, 1	AI810627	1	0.6	4.6
57347_at			AA725801	0	0.8	4.6
1827_s_at			M13929	0.9	0	4.6
64832_at			AA113447	1	0.7	4.6
41113_at		KIAA0557 protein	AI871396	1	0.4	4.5
39080_at		KDEL (Lys-Asp-Glu-Leu) endoplasmic reticulum protein retention receptor 2	M88458	0	0.3	4.5
35527_at		calcium channel, voltage-dependent, alpha 2/delta subunit 1	M76559	0.5	0.4	4.5

Figure 14S

34299_at	zinc finger protein 278	ZNF278	AL096880	1.7	2.9	4.5
35346_at	FEM-1 (C.elegans) homolog b	FEM1B	AB007856	0.8	1	4.5
32439_at	ATPase, H+/K+ exchanging, beta polypeptide	ATP4B	M75110	0.7	0.7	4.5
51367_at			A1189446	0.2	0.8	4.5
80256_at			A1167159	0	0.2	4.5
58156_i_at	peptidylprolyl isomerase A (cyclophilin A)	PPIA	R38630	0.8	0.3	4.5
44977_g_at			A1972721	0	0.1	4.5
45739_at	hypothetical protein FLJ11210		AA056538	0	0	4.5
58618_at			A1741373	0.8	0.4	4.5
46812_at			N78361	0.1	0.9	4.5
55879_at			Z40682	0.3	0.9	4.5
41856_at			AL049370	0.9	0.3	4.4
37887_at	protein kinase Chk2		AF086904	0	0	4.4
40792_s_at	triple functional domain (PTPRF interacting)	TRIO	AF091395	1	0.4	4.4
33687_at			AL049782	0.7	0.2	4.4
32739_at	KIAA1117 protein	CNTN2	AA001791	0.8	0.7	4.4
36276_at	contactin 2 (axonal)		X67734	0.5	0.4	4.4
36641_at	capping protein (actin filament) muscle Z-line, alpha 2	CAPZA2	U03851	0.4	0.7	4.4
46774_i_at			A1888795	0.3	0.2	4.4
66443_at			AA775536	0.3	0	4.4
61992_i_at			A1147067	0	0.4	4.4
84235_at	pre-B-cell leukemia transcription factor 1	PBX1	AW005924	0.6	0	4.4
63104_at			H98113	0	0.8	4.4
84699_at			N54951	0	0	4.4
72963_r_at			A1950023	0.4	0	4.4
84155_at			A1935521	0.8	0	4.4
64236_at	wingless-type MMTV integration site family, member 4	WNT4	A1634496	0.5	0.4	4.4
45613_at			AA975060	0.1	0.8	4.4
65907_r_at			A1421880	0.6	0.6	4.4
42033_at			AA243840	0.9	0.5	4.4
38285_at	crystallin, mu	CRYM	AF039397	1.1	0.6	4.4
32407_f_at			U92818	0.9	0.7	4.4
86023_at			A1478597	0	0.2	4.4
89349_at			R08903	0	0.2	4.4
38973_at	KIAA1020 protein		AB028943	1	0.8	4.3

Figure 14T

1804_at	kallikrein 3, (prostate specific antigen)	KLK3	X07730	1	0.7	4.3
1770_at	interferon, alpha 2	IFNA2	J00207	0.9	0	4.3
32140_at	sortilin-related receptor, L(DLR class) A repeats-containing	SORL1	Y08110	1.1	0.7	4.3
35987_g_at	member of MYST family histone acetyl transferases, homolog of Drosophila MOF		AL050395	0	0	4.3
36755_s_at	interleukin 5 receptor, alpha	IL5RA	M75914	0.6	0.5	4.3
84746_at			AA278620	0	0.6	4.3
64281_s_at	upregulated by 1,25-dihydroxyvitamin D-3		AW026672	0	0	4.3
86380_at			AI718112	1	0.8	4.3
67687_at	high-mobility group (nonhistone chromosomal) protein 1	HMG1	AI191227	0	0	4.3
46366_at			T79183	0.7	0.9	4.3
46885_at			AI822096	0.3	0.4	4.3
87058_at			T90289	0	0.9	4.3
35940_at	POU domain, class 4, transcription factor 1	POU4F1	X64624	0.2	0.4	4.2
38405_at	fragile X mental retardation, autosomal homolog 1	FXR1	U25165	0.7	0.9	4.2
75863_at			AL042660	0	0	4.2
61255_at			N36377	0.1	0.1	4.2
85760_r_at			AI680362	0.6	0.8	4.2
48172_at			R84376	0	0.8	4.2
79179_at			AI659829	0	0	4.2
69416_at			AI693716	0.8	0.2	4.2
80476_at			T16443	0.7	0.9	4.2
61963_at	vanilloid receptor-related osmotically activated channel; OTRPC4 protein		AI674479	0.7	0.6	4.2
58397_r_at			AA976373	0.6	0.5	4.2
63916_r_at			T50788	0.3	0.4	4.2
52446_at			AI281933	0.6	0.6	4.2
61704_i_at	hypothetical protein FLJ20036		AW006276	0.7	0.3	4.2
60581_at			AA100715	0.5	0.8	4.2
47053_at			AI768650	0	0	4.2
56735_at			AA262730	0.5	0.6	4.2
38694_at	KIAA0738 gene product		AB018281	0	0.6	4.1
35699_at	budding uninhibited by benzimidazoles 1 (yeast homolog), beta	BUB1B	AF053306	2.1	1	4.1
36075_at	putative GTP-binding protein similar to RAY/RAB1C		AL037167	0.3	0.1	4.1
36710_at	cathelicidin antimicrobial peptide	CAMP	Z38026	0.8	0.3	4.1

Figure 14U

40010_s_at	fragile X mental retardation 2	FMR2	L76569	0.4	1.1	4.1
76657_at			AA058578	0.5	0	4.1
60115_at	hypothetical protein FLJ20533		H23252	0.4	0.4	4.1
91176_at			AI792991	0.6	0.5	4.1
81074_at			AA169554	0.6	0	4.1
59487_at			R70662	0.9	0.9	4.1
54831_r_at			AA147881	0.2	0.7	4.1
71729_at			W90100	0.4	0.1	4.1
48601_at			AI525627	0.9	0.9	4.1
67812_at			AI018657	0.7	0.8	4.1
70089_at			AI765926	0.5	0.3	4.1
70352_at			AI803691	0	0.9	4.1
70831_at			AI797620	0.3	0.2	4.1
74948_at	nuclear pore complex interacting protein	NPIP	AI991242	0.9	0	4.1
77228_at			AI300953	0	0	4.1
66178_at			AA410395	0.5	0.6	4.1
65566_at	etoposide-induced mRNA		AI741005	0	0.6	4.1
51226_at			N53536	0.8	1.5	4.1
52072_at	hypothetical protein FLJ10718		AA873182	5.1	0.8	4.1
91156_at			AI219949	0.2	0.4	4.1
39016_r_at	keratin 6A	KRT6A	L42611	0.1	0.1	4
38535_at	distal-less homeobox 4	DLX4	U73328	0.8	0.8	4
1696_at	polymerase (DNA directed), beta	POLB	D29013	1.3	1.8	4
35762_at	KIAA0483 protein		AB007952	0	0.5	4
646_s_at	CDC-like kinase 2	CLK2	L29218	2.7	0.7	4
36155_at	KIAA0275 gene product		D87465	1.2	1.1	4
56847_at			AI911070	0	0	4
58206_at			W63702	0	0.3	4
56460_at			AA809379	0	0	4
59179_at			T89693	0	0	4
80872_at			AA533574	0	0.2	4
81565_at			W85907	0.7	0.4	4
60502_at			AI743953	0	0	4
71619_at			AI936553	0.9	0.9	4
61310_at	hypothetical protein FLJ10335		AL036075	0.4	0.4	4

Figure 14V

78729_at				AI939307	0.6	0.3	4
49179_at	KIAA0493 protein			AI675152	0.7	0.6	4
75607_at				AA778921	0.2	0.5	4
34539_at	olfactory receptor, family 7, subfamily E, member 12 pseudogene	OR7E12P		AF065854	0.8	0.5	3.9
39784_at				U26032	0	0.4	3.9
39459_at	ribosomal protein S13	RPS13		W28765	0.9	1	3.9
39290_f_at	PAI-1 mRNA-binding protein			W28257	0.9	0	3.9
33327_at	chromosome 11 open reading frame 9	C11ORF9		AB023171	0.1	0.7	3.9
40997_at	mitogen-activated protein kinase kinase 12	MAP3K12		AI660963	0.7	0.5	3.9
36821_at				AL050367	0	0	3.9
36754_at	adenylate cyclase activating polypeptide 1 (pituitary)	ADCYAP1		X60435	0.4	0.3	3.9
36203_at	ornithine decarboxylase 1	ODC1		X16277	1	0.3	3.9
508_at	suppressor of Ty (S.cerevisiae) 4 homolog 1	SUPT4H1		U43923	0	0.3	3.9
56434_at				AA579598	0.3	0.8	3.9
90278_at				AA709455	0.9	0.7	3.9
54777_at				H61899	0.1	0	3.9
79335_at				AA706315	0	0	3.9
43338_at				R44338	0	0.4	3.9
51868_r_at				AA829939	0	0.3	3.9
45457_at				R60981	0.2	0.1	3.9
57114_at				W87368	0.2	1	3.9
84562_r_at				W44959	1	0.5	3.9
44528_at	hypothetical protein FLJ20094			AI480121	0.7	0.7	3.9
53185_at	collapsin response mediator protein-5; CRMP3-associated molecule			AI879238	0.9	0.7	3.9
37475_at	DKFZP434J046 protein			AC004144	0.5	0.9	3.8
40634_at	nucleosome assembly protein 1-like 1	NAP1L1		M86667	0.4	0.5	3.8
33309_at				AA521060	0.1	0.3	3.8
41440_at	FabG (beta-ketoacyl-[acyl-carrier-protein] reductase, E coli) like	FABGL		D82061	1.7	1.6	3.8
799_at	cyclin-dependent kinase 5, regulatory subunit 1 (p35)	CDK5R1		X80343	1	0.3	3.8
36281_at				M96739	0.4	0.2	3.8
40030_at	protein kinase, Y-linked	PRKY		Y15801	0	0.8	3.8
38764_at	helicase-moi			AF007142	0.2	0	3.8
76724_r_at				H38425	0.2	1	3.8

Figure 14W

78106_at			AI634844	0.1	0	3.8
90840_at			W88900	0.8	0.3	3.8
79388_at			AI128130	0.9	0.8	3.8
84492_at			AI288213	0	0.9	3.8
44832_at			R26252	0	0	3.8
75162_r_at	erythrocyte membrane protein band 4.9 (dematin)	EPB49	AI673236	1	0.5	3.8
58734_at			N48694	0	0	3.8
46911_at			AI381767	1	0.3	3.8
47106_at	hypothetical protein MGC3169		AA196034	0.2	0.4	3.8
45488_s_at			AI525989	0.9	0.5	3.8
887_at	growth differentiation factor 1	GDF1	M62302	0.8	0.3	3.8
55481_i_at			R54660	0.3	0.2	3.8
88335_r_at			AA126277	0.9	0.6	3.8
34902_at	KIAA0492 protein		AB007961	0.1	0	3.7
41282_s_at	peroxisome biogenesis factor 10	PEX10	AA194159	0.5	0.4	3.7
35138_at	A kinase (PRKA) anchor protein 8	AKAP8	Y11997	0	0.1	3.7
40595_at	Treacher Collins-Franceschetti syndrome 1	TCOF1	AI345337	1.5	0.3	3.7
36580_at	hypothetical protein FLJ13910		AL050139	0	0.7	3.7
38126_at	biglycan	BGN	J04599	0	0	3.7
35028_at	gamma-aminobutyric acid (GABA) A receptor, beta 1	GABRB1	X14767	0	0	3.7
43059_at	KIAA0955 protein		AI040349	0	0	3.7
61471_at			N66393	0	0	3.7
75557_at			AI446519	0.7	0.7	3.7
52763_at			AI277429	1	0.7	3.7
45240_at	stromal cell protein		N93439	0.5	0.6	3.7
75254_at	matrilin 2	MATN2	AI972357	0.8	0	3.7
71309_at			AI919139	0.3	0.9	3.7
71504_at			AA811476	1.8	1.2	3.7
59632_at			AI991777	0.1	1	3.7
72080_at			N99035	0	0.3	3.7
64262_at			R39949	0.9	0.6	3.7
50933_at	Alport syndrome, mental retardation, midface hypoplasia and elliptocytosis chromosomal region, gene 1	AMMECR1	AL037664	0	0.1	3.7
57757_r_at			AA425516	0	0	3.7
80081_at	ribosomal protein L4	RPL4	AI721208	0.5	0.5	3.7

Figure 14X

55971_at	KIAA1323 protein		AI263115	0	0	3.7
54477_at			AI193993	0	1.1	3.7
68127_at			AA308556	0	0	3.7
37459_at	collagen, type VIII, alpha 1	COL8A1	X57527	0.2	0	3.6
39694_at	hypothetical protein MGC5508		W27517	0	0.5	3.6
35325_at	GTPase Rab14		AF052113	0.5	0.8	3.6
1083_s_at	mucin 1, transmembrane	MUC1	M35093	0.1	0.2	3.6
57379_at			AA865085	1	0.6	3.6
50282_at	hypothetical protein FLJ10975		AA522810	0	0.2	3.6
88783_at			AI961262	0.9	0.8	3.6
53745_at	LCHN protein		AI345949	0.2	0.7	3.6
74529_at			AI056998	0.4	0.2	3.6
82705_at			AA464792	0.5	0.9	3.6
85376_at			AL042726	0.3	0	3.6
56275_at			AI631775	0.5	1	3.6
71407_at			AI651482	0.9	0.2	3.6
69003_at			AA602488	0.4	0.7	3.6
51766_at			AI472196	0	0.2	3.6
46130_at			AL042817	0	0	3.6
47384_at			AI806045	0	0.5	3.6
42000_r_at			T40827	0.3	0.1	3.6
46050_r_at			AL120288	0.2	0.1	3.6
39361_f_at	transmembrane 4 superfamily member 6	TM4SF6	AF043906	0	0	3.5
41712_at	KIAA0560 gene product		AB011132	0	0.2	3.5
35441_at	GLE1 (yeast homolog)-like, RNA export mediator	GLE1L	AF058922	0	1	3.5
33280_r_at	SA (rat hypertension-associated) homolog	SAH	X80062	0.5	0.1	3.5
32180_s_at	gene with multiple splice variants near HD locus on 4p16.3		AB000461	1	0.9	3.5
34305_at	poly(rC)-binding protein 1	PCBP1	Z29505	0.7	0.6	3.5
772_at	v-crk avian sarcoma virus CT10 oncogene homolog	CRK	D10656	0	1.2	3.5
34607_at	inducible T-cell co-stimulator	ICOS	AB023135	0.6	0.5	3.5
40646_at	chemokine (C-X3-C) receptor 1	CX3CR1	U20350	0	0.8	3.5
39137_at	nuclear factor related to kappa B binding protein	NFRKB	X80878	0.8	0	3.5
64361_s_at	ubiquitin-conjugating enzyme E2L 3	UBE2L3	AW006924	0	0	3.5
56431_at	hematological and neurological expressed 1	HN1	N90863	0	0	3.5
75843_at			AA808011	0.4	0.1	3.5

Figure 14Y

47199_at			AL119929	2.3	2	3.5
59910_at			AA908954	0.6	0	3.5
82355_at			AI358881	0.7	0	3.5
59507_at			AI638235	0	0.6	3.5
44950_at			N92407	1	0.5	3.5
53534_at			N63469	0.7	0.8	3.5
83721_at			W56091	0.7	0.9	3.5
48728_r_at			W74754	0.3	0.1	3.5
42112_at			AA398813	0.1	0.1	3.5
52351_at			3053308	0.1	0.5	3.5
1158_s_at	calmodulin 3 (phosphorylase kinase, delta)	CALM3	J04046	0.4	0.1	3.4
34195_at			AL121073	0.4	1	3.4
34012_at	keratin, hair, acidic, 4	KRTHA4	Y16790	0.4	0	3.4
38122_at	solute carrier family 23 (nucleobase transporters), member 1	SLC23A1	D87075	0	0	3.4
56490_at			AA682764	0.4	0.5	3.4
84624_at			AA505694	0.6	0.5	3.4
79835_at			AA203646	0	0.1	3.4
43025_r_at			AI821404	0.7	0.9	3.4
47539_at			AI807478	0.5	0.6	3.4
44516_at			AI809013	1	0.2	3.4
44626_g_at			R78655	0.9	0.9	3.4
56861_at	Rap2 interacting protein 8		AI197870	1	0.4	3.4
31535_i_at			W27858	0.4	0	3.4
37150_at	Kelch motif containing protein		AB026190	1.2	0	3.3
39550_at	KIAA0584 protein		AB011156	0	0	3.3
35533_f_at	killer cell lectin-like receptor subfamily C, member 4	KLRC4	AJ001683	0	0	3.3
34873_at	nebulette		Y16241	0.3	0	3.3
37710_at	MADS box transcription enhancer factor 2, polypeptide C (myocyte enhancer factor 2C)	MEF2C	L08895	0.8	0	3.3
38470_i_at	amyloid beta precursor protein (cytoplasmic tail)-binding protein 2	APPBP2	D86981	0.7	0.3	3.3
37928_at	nuclear transcription factor Y, beta	NFYB	AA621555	0.3	0	3.3
33191_at	hypothetical protein FLJ10512		AW051579	0.8	0	3.3
42369_at			R43543	0	0.3	3.3
57053_s_at	homolog of rat nadrin		AI189504	0.1	0.8	3.3

Figure 14Z

79437_at			AI138554	0.4	0	3.3
52635_at			AI820880	0.1	0	3.3
59058_at			AA404351	0	0	3.3
87852_at			AI479665	0.7	0	3.3
62820_at			AW002871	0.4	0.6	3.3
61731_at			W69427	0.8	0	3.3
50440_at		hypothetical protein FLJ23342	AA393461	0.9	0.6	3.3
1415_at			D26561	0.4	0.8	3.3
61252_f_at			H60379	0.6	0	3.3
52825_f_at			AI251752	0.8	0	3.3
67654_r_at			AA435672	0	0.6	3.3
82498_at			AI479524	0	0.2	3.3
31417_at		hypocretin (orexin) neuropeptide precursor	AF041240	0	0.8	3.2
39450_s_at		iduronate 2-sulfatase (Hunter syndrome)	AF050145	0.1	0.8	3.2
40927_at		solute carrier family 6 (neurotransmitter transporter, creatine), member 8	W26324	0.8	0.1	3.2
33794_g_at		transcription factor 20 (AR1)	U19345	0.9	0.3	3.2
33017_at		glycosylphosphatidylinositol specific phospholipase D1	AI040379	0.8	0	3.2
34778_at		NeuAc-alpha-2,3-Gal-beta-1,3-GalNAc-alpha-2, 6-sialyltransferase	AA418080	0.4	0.2	3.2
33640_at		alpha2,6-sialyltransferase	Y14768	0.2	0	3.2
33109_f_at		allograft inflammatory factor 1	L07335	1	0.7	3.2
41034_s_at		SRX (sex determining region Y)-box 2	U92315	0.6	0.9	3.2
77799_at		sulfoltransferase family, cytosolic, 2B, member 1	AI632915	1	0.7	3.2
47463_at		eukaryotic translation initiation factor 2C, 1	AA453723	0	0	3.2
84063_at			AI417354	2.1	1	3.2
82457_at			AI798172	0	0	3.2
43314_at			AA009521	1	1.6	3.2
50900_at			AI991581	0.5	0.9	3.2
61032_at			AI122634	0.3	0.4	3.2
44616_r_at			AA057585	0.5	0	3.2
52868_g_at		hypothetical protein FLJ20279	AA039793	0	0	3.2
727_at				1	0.7	3.2
67665_f_at			R02298	0.5	0	3.2
41466_s_at		zinc finger protein 148 (pH2-52)	L04282	0	0.2	3.1

Figure 14AA

878_s_at	prolactin	PRL	M29386	0	0	3.1
34089_at	KIAA1030 protein		AB028953	0.9	0	3.1
31655_at			AL031737	1	0.5	3.1
31699_at	phosphoinositide-3-kinase, catalytic, beta polypeptide	PIK3CB	S67334	0.4	0	3.1
35566_f_at	immunoglobulin heavy constant mu	IGHM	AF015128	1	0.9	3.1
1794_at	cyclin D3	CCND3	M92287	0	0	3.1
38364_at	BCE-1 protein		AF068197	0	0.9	3.1
65433_at			T86971	0	0.1	3.1
63407_at	cell division cycle 27	CDC27	AA142869	0.9	0.8	3.1
48231_at			H08161	0.8	0.8	3.1
86639_at			AI697562	0.3	0.5	3.1
43869_at			R49539	0.9	0	3.1
90775_at			AI821755	0.9	0	3.1
84545_at			AA601040	0.9	0.5	3.1
64121_at	glioma-amplified sequence-41		Z78373	0	1	3.1
45073_at			N69578	0.1	1.7	3.1
47346_at			AI818064	0	0.3	3.1
55007_at			AA521088	0.3	1	3.1
48358_at			AA760689	0.9	1	3.1
1180_g_at				0	0.9	3.1
88334_f_at			AA586897	0.7	0	3.1
39899_at			AC005525	0.7	0.6	3
709_at	tubulin, beta polypeptide	TUBB	J00314	0.4	0.5	3
32292_at	collectin sub-family member 10 (C-type lectin)	COLEC10	AB002631	0.6	0.3	3
37488_at	farnesyltransferase, CAAX box, beta	FNTB	L00635	0.7	0.4	3
34796_at	translocating chain-associating membrane protein		X63679	0	0.6	3
37402_at	ribonuclease, RNase A family, 1 (pancreatic)	RNASE1	D26129	0.6	0.3	3
38343_at	KIAA0328 protein		AB002326	0.8	1	3
58055_at			AI023433	0.5	1	3
48546_s_at	eukaryotic translation elongation factor 1 alpha 1	EEF1A1	N30008	0.6	1	3
45680_at	KIAA1474 protein		AI741882	0	0	3
45155_at	claudin 15	CLDN15	AI433892	1	0.9	3
92097_at			AI935368	0.2	0.5	3
72751_at			N67652	0.7	0.3	3

Figure 14BB

affyID	Gene Name	Gene Symbol	GBA	Score_Ecoli	Score_Candida	Score_Influenza
35061_at	small inducible cytokine subfamily B (Cys-X-Cys), member 11	SCYB11	AF030514	655.1	844.8	729.7
563_at	interleukin 12B (natural killer cell stimulatory factor 2, cytotoxic lymphocyte maturation factor 2, p40)	IL12B	M65290	2449.1	761.6	1.2
425_at	interferon, alpha-inducible protein 27	IFI27	X67325	565.9	754	421
37219_at	monokine induced by gamma interferon	MIG	X72755	261.6	704.2	182.7
38549_at			AF026941	484.1	612.9	1149.5
431_at	small inducible cytokine subfamily B (Cys-X-Cys), member 10	SCYB10	X02530	152	390	545.7
37944_at	GTP cyclohydrolase 1 (dopa-responsive dystonia)	GCH1	U19523	505.4	330.9	169
35822_at	B-factor, properdin	BF	L15702	1635.7	287.1	17.3
1520_s_at	endothelin 1	EDN1	J05008	4444	283.5	113.7
48864_at	interferon, alpha-inducible protein 27	IFI27	A1991845	216.4	274.9	141.4
39959_at	gamma-aminobutyric acid (GABA) B receptor, 1	GABBR1	AL031983	161.4	263.9	33.7
33304_at	interferon stimulated gene (20kD)	ISG20	U88964	327.2	200	103.7
52615_at			AA948319	425	188.4	94.9
1715_at	tumor necrosis factor (ligand) superfamily, member 10	TNFSF10	U37518	66.8	179.7	463.3
70458_i_at			A1654525	120.3	174.3	127.2
62330_at			A1075407	185.9	169.1	121.9
51972_at	ubiquitin specific protease 18	USP18	AA143794	114.4	158.3	218
38584_at	interferon-induced protein with tetratricopeptide repeats 4	IFIT4	AF026939	179.4	153.2	148.6
36776_at	lymphocyte-activation gene 3	LAG3	X51985	13.2	148.7	69.7
61371_at	cyclin-E binding protein 1		AA555023	204	148	479.5
909_g_at	interferon-induced protein with tetratricopeptide repeats 2	IFIT2	M14660	320.5	145.5	573.3
675_at	interferon induced transmembrane protein 1 (9-27)	IFITM1	J04164	112.7	140.9	173.6
84893_at			A1446168	83.4	119.5	160.2
40323_at	CD38 antigen (p45)	CD38	D84276	160.1	102.8	18.6
38299_at	interleukin 6 (interferon, beta 2)	IL6	X04430	1566.4	101.3	227.7
1097_s_at	chemokine (C-C motif) receptor 7	CCR7	L31584	276.6	100.6	88.3
41677_at	interleukin 15 receptor, alpha	IL15RA	AF035279	278	99.5	111.4
40159_r_at	neutrophil cytosolic factor 1 (47kD, chronic granulomatous disease, autosomal 1)	NCF1	M55067	160.7	91	140.3
1403_s_at	small inducible cytokine A5 (RANTES)	SCYA5	M21121	232.1	90.6	200.3
70460_f_at			A1654525	48.9	90.2	75.7
1107_s_at	interferon-stimulated protein, 15 kDa		M13755	159	86.3	213.8

Figure 15A

53508_at				AI247103	124.7	84.6	27.2
37014_at	myxovirus (influenza) resistance 1, homolog of murine (interferon-inducible protein p78)	MX1		M33882	53.6	82.5	69.2
71174_r_at	KIAA1682 protein			AI921158	67.8	81.9	115.6
44513_at				AI760613	68.8	81.5	36.6
51546_at				AI498361	113.1	77.9	73.9
91313_at				AA131041	84	77.5	157.1
74689_at	lymphocyte-activation gene 3	LAG3		AW003135	7.7	77.5	26.1
71839_at				AA251131	111.4	76.5	82.8
89882_at	interferon induced transmembrane protein 1 (9-27)	IFITM1		AI656093	38.3	76.3	71.5
74566_at	insulin-like growth factor-binding protein 4	IGFBP4		AI440424	13.7	75.4	14.1
45164_at	19A24 protein			W74027	189.7	74.5	32.5
41745_at	interferon induced transmembrane protein 3 (1-8U)	IFITM3		X57352	49.6	74	69.9
65867_at	FXYD domain-containing ion transport regulator 6	FXYD6		AL043089	103.8	73.3	28.2
1737_s_at	insulin-like growth factor-binding protein 4	IGFBP4		M62403	26.2	72	18.6
38389_at	2',5'-oligoadenylate synthetase 1 (40.46 kD)	OAS1		X04371	36.5	71.8	35
32814_at	interferon-induced protein with tetratricopeptide repeats 1	IFIT1		M24594	50.5	70.1	76.9
32775_r_at	phospholipid scramblase 1	PLSCR1		AB006746	81.8	68.4	32.6
39775_at	serine (or cysteine) proteinase inhibitor, clade G (C1 inhibitor), member 1	SERPING1		X54486	41.5	67.8	17.1
41045_at	secreted and transmembrane 1	SECTM1		U77643	37.2	67.4	10.4
2002_s_at	BCL2-related protein A1	BCL2A1		U27467	547.8	65.2	70.5
89899_at	myxovirus (influenza) resistance 2, homolog of murine	MX2		AW001846	54	64.3	70.9
33789_at	small inducible cytokine A5 (RANTES)	SCYA5		AF088219	191.5	64.1	1.2
37823_at	small inducible cytokine subfamily A (Cys-Cys), member 8 (monocyte chemotactic protein 2)	SCYA8		Y16645	17.3	63.9	47.1
88714_at				AA873008	94.5	63.3	12.9
59777_at				AA766831	98.3	63.3	41.1
34666_at	superoxide dismutase 2, mitochondrial	SOD2		X07834	141.3	63.1	42.3
59215_at	RNA helicase			AI807018	151.1	63	36.2
36296_at	lymphotoxin alpha (TNF superfamily, member 1)	LTA		D12614	154	62.8	28.5
879_at	myxovirus (influenza) resistance 2, homolog of murine	MX2		M30818	61.1	61	69
36927_at	hypothetical protein, expressed in osteoblast			AB000115	53.4	60.4	83.5
77706_at				AA641972	493	59	7.8
1021_at	interferon, gamma	IFNG		J00219	289.9	58.8	36.2

Figure 15B

71250_at				AI984837	54.1	57.5	0
36116_at	thyroid hormone receptor interactor 10	TRIP10		AJ000414	215.9	56.7	7
63335_at	sorting nexin 10			AI285531	127.4	55.8	74.2
90956_at				AA932068	99.3	55.6	0.8
38570_at	major histocompatibility complex, class II, DO beta	HLA-DOB		X03066	84.2	55.5	30.1
84909_at				AA902134	29.7	55.3	39.5
33236_at	retinoic acid receptor responder (tazarotene induced) 3	RARRES3		AF060228	9.2	55.2	24.8
46659_at				AI742057	87.7	54.8	56
38388_at	2',5'-oligoadenylate synthetase 1 (40-46 kD)	OAS1		M11810	38.7	54	103.7
50827_at				AW001213	109.2	52.4	6.6
49148_at				AA947472	182.2	52	423.6
33705_at	phosphodiesterase 4B, cAMP-specific (dunce (Drosophila)-homolog phosphodiesterase E4)	PDE4B		L20971	232.3	51.8	92.7
87102_at				AI823649	62	51.5	48.7
47517_at				AA639645	67.1	51.4	3.3
37641_at	interferon-induced, hepatitis C-associated microtubular aggregate protein (44kD)			D28915	68.5	51.1	43.6
1527_s_at				U50527	82.5	50.7	14.2
37168_at	similar to lysosome-associated membrane glycoprotein			AB013924	90.9	50.6	16.1
57761_at				AI313253	161	50.6	33.7
35735_at	guanylate binding protein 1, interferon-inducible, 67kD	GBP1		M55542	72.1	49.4	49.8
38787_at	promyelocytic leukemia	PML		X63131	17.9	49.4	18.9
63769_at				AI749656	67.8	49	14.3
1717_s_at	baculoviral IAP repeat-containing 3	BIRC3		U45878	63.1	47.7	20.2
1852_at	tumor necrosis factor (TNF superfamily, member 2)	TNF		X02910	201.2	47.5	57
40447_at	lipin 2	LPIN2		D87436	20.2	47.2	18.2
36575_at	regulator of G-protein signalling 1	RGS1		S59049	31	46.9	16.7
40679_at	solute carrier family 6 (neurotransmitter transporter, betaine/GABA), member 12	SLC6A12		U27699	80.9	46.2	10.4
85586_at	melanoma antigen, family D, 1	MAGED1		AI336885	21.4	46.1	22.3
41744_at	tumor necrosis factor alpha-inducible cellular protein containing leucine zipper domains; Huntingtin interacting protein L; transcription factor IIIA-interacting protein			AF070533	26	45.9	21.1
38121_at	tryptophanyl-tRNA synthetase	WARS		X59892	58.2	45.4	45.6
89884_at	metallothionein 2A	MT2A		AI991852	138	45.1	82.3

Figure 15C

39710_at	P311 protein		U30521	6.6	35	8.7
46133_at			W60263	34.8	34.9	12.6
1761_at	platelet-derived growth factor receptor-like	PDGFR	D37965	11.9	34.7	14.4
39119_s_at	natural killer cell transcript 4		AAG31972	22.5	34.2	10.5
62130_at	28kD interferon responsive protein		AA651720	37.5	34	40
908_at	interferon-induced protein with tetratricopeptide repeats 2	IFIT2	M14660	22.9	34	384.9
37220_at	Fc fragment of IgG, high affinity Ia, receptor for (CD64)	FCGR1A	M63835	0.5	33.7	2.3
37216_at	KIAA0963 protein		AB023180	75.3	33.5	1.2
63927_f_at			AI963642	38.5	33.4	11.9
39271_at			AA461365	7.6	33.2	1.5
40153_at	ATP-binding cassette, sub-family B (MDR/TAP), member 2	RAB4B	X57522	34.3	33	40.6
1369_s_at	interleukin 8	ABC2	M28130	2300.5	32.6	122.1
45237_at	superoxide dismutase 2, mitochondrial	IL8	AA142976	84.4	32.6	17.7
61818_at		SOD2	AI970348	265	32.3	2.5
69876_at			AI825713	36.1	32.1	60.3
37187_at	GRO2 oncogene	GRO2	M36820	785	32	26.1
60896_at	integrin, beta 8	ITGB8	AI634852	137.5	31.7	10.7
57699_at			AA133285	12.1	31.2	65.3
48018_at			AA700227	19.1	31.1	9.3
32114_s_at	adenosine A2a receptor	ADORA2A	S46950	137.9	31	27.9
33823_at	CD36 antigen (collagen type I receptor, thrombospondin receptor)-like 2 (lysosomal integral membrane protein II)	CD36L2	D12676	16.8	30.9	9.5
1370_at	interleukin 7 receptor	IL7R	M29696	102.3	30.8	25.3
39402_at	interleukin 1, beta	IL1B	M15330	468.5	30.5	12.2
35669_at	KIAA0633 protein		AB014533	3.2	30.4	24.1
89202_at			AI347361	28.3	30.4	9.1
64450_at	2'-5'-oligoadenylate synthetase 3	OAS3	AA135525	25.2	30.3	10.8
54823_at			AI651594	35.1	30	5.8
70097_at	indoleamine-pyrrole 2,3 dioxygenase	INDO	AI302268	99.5	29.8	39.9
50124_at			AA580047	37.3	29.6	43.2
33364_at	similar to rat myomegalin		U51694	156	28.9	2.9
61940_at	hypothetical protein FLJ21343		W28128	12	28.9	20.7
87002_i_at			AI270476	14.8	28.7	63.7
62214_at	hypothetical protein FLJ11354		AI583960	35.5	28.7	23.8
44126_at			W28501	73.5	28.6	16.2

Figure 15E

40385_at	small inducible cytokine subfamily A (Cys-Cys), member 20	SCYA20	U64197	943.4	28.3	53.8
79210_at			W58117	2.6	28.1	0.2
41742_s_at	tumor necrosis factor alpha-inducible cellular protein containing leucine zipper domains; Huntingtin interacting protein L; transcription factor IIIA-interacting protein		AF061034	24.3	28.1	8.7
39263_at	2'-5'oligoadenylate synthetase 2	OAS2	M87434	26.7	28	
91636_at	KIAA0580 protein		AB011152	886.7	27.7	34.8
39224_at	2'-5'oligoadenylate synthetase 2	OAS2	AB011152	71.4	27.4	7
90662_at	JM5 protein		AB011152	25.2	27.3	18
33350_s_at	TNF receptor-associated factor 1	TRAF1	Z78315	37.4	27.2	19
90942_at			AF117829	97.1	27	1
55776_at	receptor-interacting serine-threonine kinase 2	RIPK2	W20082	16.8	26.9	0.5
41384_at			AF117829	46.7	26.8	17.7
80045_at			AF117829	29.1	26.8	104.9
1069_at	prostaglandin-endoperoxide synthase 2 (prostaglandin G/H synthase and cyclooxygenase)	PTGS2	U04636	495.3	26.7	12.8
39114_at	decidual protein induced by progesterone					53.1
35015_at	CD80 antigen (CD28 antigen ligand 1, B7.1 antigen)	CD80	AB022718	68.6	26.6	
82359_at			M27533	61	26.5	28.6
59478_at			AB022718	15.7	26.5	15.7
60903_s_at	T-cell lymphoma invasion and metastasis 2	TIAM2	AA741304	10.6	26.4	44.6
34459_at			AB022718	41.6	25.5	11.9
36067_at	small inducible cytokine subfamily A (Cys-Cys), member 19	SCYA19	U50529	11.7	25.2	6.8
32026_s_at	PDZ domain containing guanine nucleotide exchange factor(GEF)1		AB000887	394.2	25.2	0.7
1842_at			AB002311	68.3	25.1	2.8
1867_at	CASP8 and FADD-like apoptosis regulator		S62138	11.6	25	16.6
41474_at	kinesin heavy chain member 2	CFLAR	AF005775	54.9	24.7	46.1
190_at	nuclear receptor subfamily 4, group A, member 3	KIF2	Y08319	13.3	24.6	5.8
59283_at	hypothetical protein FLJ20035	NR4A3	U12767	29.1	24.6	7.3
72660_g_at			AL042790	37.5	24.6	25.8
49117_at	interferon-induced protein 75, 52kD		AA776412	56.8	24.4	11.4
50296_i_at	apolipoprotein L, 3	IFI75	AF1743445	26.5	24.4	33.3
50361_at		APOL3	T63428	20	24.4	71
35143_at	KIAA1031 protein		AF1658662	17.1	24.2	12.1
58941_at	phosphatidylserine-specific phospholipase A1alpha		AF138605	50.6	24.1	11.4
			AF165967	28.1	24.1	4.6
						11.5

Figure 15F

41872_at	deafness, autosomal dominant 5	DFNA5	AF073308	27.4	24	1.3
58918_at			AA210892	176.1	24	34.8
32700_at	guanylate binding protein 2, interferon-inducible	GBP2	M55543	55.5	23.9	6.1
32529_at	transmembrane protein (63kD), endoplasmic reticulum/Golgi intermediate compartment		X69910	21.7	23.9	6.8
39070_at	singed (Drosophila)-like (sea urchin fascin homolog like)	SNL	U03057	36.9	23.6	31
53236_at	carbohydrate (N-acetylglucosamine 6-O) sulfotransferase 7	CHST7	N64774	46.8	23.4	1.6
62528_at	hypothetical protein FLJ20073		AA741307	51.4	23.4	10.3
54594_at	KIAA0668 protein		AA009571	78.6	23.3	31.3
64530_at			AA723927	17.6	23.2	15.8
44289_at	NY-REN.18 antigen		AA587445	59.8	23.1	1.9
47484_at			H17489	32.5	22.9	8.8
62804_at			AA621888	0.4	22.9	0.2
52062_f_at			H98105	16.3	22.9	6.4
32595_at	G-rich RNA sequence binding factor 1	GRSF1	U07231	7.8	22.7	16.5
31623_f_at	metallothionein 1A (functional)	MT1A	K01383	160.3	22.3	49.7
36825_at	stimulated trans-acting factor (50 kDa)		X82200	18.3	22.2	11.5
63758_at	promyelocytic leukemia	PML	A1270297	5.6	22	6.3
36227_at			AF043129	63.4	21.8	18.5
75510_at			A1821517	0	21.7	3.2
38006_at	CD48 antigen (B-cell membrane protein)	CD48	M37766	15.4	21.6	9.8
51015_at			A1655467	64.4	21.6	4.4
87497_f_at			A1832016	27	21.1	7.7
40074_at	methylene tetrahydrofolate dehydrogenase (NAD+ dependent), methenyltetrahydrofolate cyclohydrolase	MTHFD2	X16396	18.4	21	0
286_at	H2A histone family, member O	H2AFO	L19779	64.6	21	57.1
49438_at			A1535730	68.4	21	15.8
55634_at			A1040355	35.9	21	3.9
52085_at			AA210833	33.1	21	30.6
38326_at	putative lymphocyte G0/G1 switch gene		M69199	64.6	20.8	6.5
44087_at	HSPC018 protein		A1279029	21.5	20.8	2.1
58605_at	DNA segment on chromosome X (unique) 9928 expressed sequence	DXS9928E	A1697939	21.5	20.6	9.4
65626_at						
34916_s_at	tumor necrosis factor receptor superfamily, member 4	TNFRSF4	AA059458	96.4	20.4	9
			S76792	35.7	20.3	7.8

Figure 15G

35463_at	homeo box (expressed in ES cells) 1	HESX1	AF059734	19.4	20.3	40.2
946_at	t-complex-associated-testis-expressed 1-like 1	TCTEL1	D50663	22.1	20.3	12.3
38788_at	promyelocytic leukemia	PML	M82827	25.1	20.3	8.8
39420_at			S62138	11.6	20.2	42.5
35985_at	A kinase (PRKA) anchor protein 2	AKAP2	AB023137	49	20.1	24.6
50746_at			AA897516	71	20.1	43.9
43500_at	hypothetical protein FLJ20693		AI674483	11.1	19.9	2.3
50210_at			AA429326	5.8	19.9	21.3
37754_at	lectin, galactoside-binding, soluble, 3 binding protein (galectin 6 binding protein)	LGALS3BP	L13210	10.5	19.8	7.3
81966_at			AI199418	31.2	19.8	24.4
91017_at	baculoviral IAP repeat-containing 3	BIRC3	U54711	19.2	19.6	3
63780_at	hypothetical protein FLJ11259		AA814195	51.4	19.6	26.4
81915_at	cytotoxic T-lymphocyte-associated protein 4	CTLA4	AI733018	0	19.5	9.8
33386_at	H1 histone family, member 0	H1FO	Z97630	25.9	19.3	1.4
74611_s_at	complement component 4A	C4A	AI983615	1.5	19.3	8.4
36600_at	proteasome (prosome, macropain) activator subunit 1 (PA28 alpha)	PSME1	L07633	15.9	19.1	6.7
40419_at	erythrocyte membrane protein band 7.2 (stomatrin)	EPB72	X85116	42.8	19	21.9
40852_at	tudor repeat associator with PCTAIRE 2		AB025254	20.7	18.9	29.9
90663_at	suppressor of Ty (S.cerevisiae) 3 homolog	SUPT3H	AA215579	11.6	18.9	2.9
75146_at			AL047300	94.9	18.8	0.5
58873_at	neuropilin 2	NRP2	AA778058	27.2	18.8	1.1
63390_at	KIAA0603 gene product		AI948512	17.5	18.7	12
1326_at	caspase 10, apoptosis-related cysteine protease	CASP10	U60519	7.2	18.6	15.9
88629_at			AI337136	42	18.5	15.7
33955_at	interleukin 12A (natural killer cell stimulatory factor 1, cytotoxic lymphocyte maturation factor 1, p35)	IL12A	M65291	250.2	18.4	14.6
43524_at	KIAA1170 protein					
1377_at	nuclear factor of kappa light polypeptide gene enhancer in B-cells 1 (p105)	NFKB1	C14031	42.1	18.4	7.2
49459_at	apolipoprotein L	APOL1	M58603	54.5	18.3	18
56922_at			AA156784	25.2	18.2	17.4
55362_at			AA211158	32.7	18.2	35.1
34439_at	absent in melanoma 2		AI653767	4.6	18.2	19.8
33106_at	nuclear receptor subfamily 1, group H, member 3	AIM2	AF024714	40.9	18.1	16
		NRIH3	U22662	13.2	18.1	18

Figure 15H

32859_at	signal transducer and activator of transcription 1, 91kD	STAT1	M97935	3.2	18	8.2
81774_at			AI683036	10.6	18	0.6
38631_at	tumor necrosis factor, alpha-induced protein 2	TNFAIP2	M92357	63.8	17.8	18.4
32606_at	brain acid-soluble protein 1	BASP1	AA135683	56.9	17.8	11
40957_at	KIAA0160 protein	KIAA0160	D63881	14.6	17.7	5
44092_at			AI862887	21.3	17.7	5.5
33055_at	cytochrome P450, subfamily VIIIB (oxysterol 7 alpha-hydroxylase), polypeptide 1	CYP7B1	AF029403	99.5	17.6	1.2
53773_at	tumor necrosis factor (ligand) superfamily, member 13b	TNFSF13B	AA682496	16.9	17.6	11.2
37407_s_at	myosin, heavy polypeptide 11, smooth muscle	MYH11	AF013570	87	17.5	7
44825_at	RAD50 (S. cerevisiae) homolog	RAD50	AA126482	27.8	17.5	30.9
37643_at	tumor necrosis factor receptor superfamily, member 6	TNFRSF6	X63717	69	17.4	30
62495_f.at			AI433785	4.4	17.4	28.1
53670_at			AA664156	4.7	17.4	9.1
73972_at			AI863446	33.1	17.3	2.3
51024_at			AI400509	13.3	17.3	7.2
38909_at	cytochrome P450, subfamily XXVIIIB (25-hydroxyvitamin D-1-alpha-hydroxylase), polypeptide 1	CYP27B1	AB005038	98.2	17.2	2.9
35937_at	MHC class I polypeptide-related sequence B	MICB	U65416	16.3	17.2	32.3
68002_at			AA503803	21.9	17.2	13.1
48964_at			AI629027	24.2	17.2	9.1
50297_f.at	apolipoprotein L, 3	APOL3		10	17.1	7.3
76364_at			AA643507	6.4	17.1	23.3
47939_at			AA102788	0	17.1	11.2
48885_at			R61847	2.1	17.1	40
106_at	runt-related transcription factor 3	RUNX3	Z35278	20.4	17	11.3
65895_at	DKFZP564A2416 protein		AL040007	13	17	1.2
47899_at			AA056755	23.8	16.9	13.4
48039_at			AI634580	15	16.9	9.4
45146_at			AI693688	25.6	16.7	0
55051_at			AA005281	14.7	16.7	12.4
1036_at	interleukin 15	IL15	U14407	64.7	16.6	22.1
62939_at			D12309	21.7	16.6	4
63984_f.at			AA535914	15	16.4	9.2
32745_at	nuclear localization signal deleted in velocardiofacial syndrome	NLVCF	AF034091	2.4	16.3	6.8

Figure 15I

35957_at	stannin	SNN	AF030196	22.2	16.3	1.1
31693_f_at	H2A histone family, member G	H2AFG	Z80776	29.8	16.1	30.5
38986_at	glucose regulated protein, 58kD	GRP58	Z49835	7.9	16.1	4.7
67401_at			AI434675	16.7	16.1	6.4
1914_at	cyclin A1	CCNA1	U66838	57.6	16	70.3
57221_g_at	proteasome (prosome, macropain) activator subunit 1 (PA28 alpha)	PSME1	AA662670	10.4	15.9	2.5
32200_at	acid phosphatase, prostate	ACPP	M24902	1.5	15.8	0.4
38287_at	proteasome (prosome, macropain) subunit, beta type, 9 (large multifunctional protease 2)	PSMB9	AA808961	15.2	15.8	3
1633_g_at	pim-2 oncogene	PIM2	U77735	32	15.7	11.4
62743_at	Wiskott-Aldrich syndrome protein interacting protein	WASPIP	AA811088	1.9	15.6	0
64930_at	FN5 protein		AA587944	22.3	15.6	3.9
1173_g_at				26.2	15.6	19.2
37416_at	ras homolog gene family, member H	ARHH	Z35227	16.4	15.5	11.5
32684_at			AF038174	13.1	15.5	2.9
74633_at	tumor necrosis factor, alpha-induced protein 2	TNFAIP2	AI986430	53.5	15.5	6.1
46266_at	serine (or cysteine) proteinase inhibitor, clade A (alpha-1 antiproteinase, antitrypsin), member 8	SERPINA8	AI208913	23.9	15.5	9.9
41147_at			AF038186	4.1	15.4	3.6
77244_at	bHLH factor Hes4		AI380603	64.7	15.4	23.6
63785_at	Mitochondrial Acyl-CoA Thioesterase		AI074605	13	15.4	3.2
52183_at			AA243659	33.5	15.4	18.8
34435_at	aquaporin 9	AQP9	AB008775	50.5	15.3	3.4
43853_at	hypothetical protein		AA618602	296.2	15.3	18.6
72862_at			AW026509	11.9	15.3	10
37294_at	B-cell translocation gene 1, anti-proliferative	BTG1	X61123	34.3	15.2	8
55001_at			R13843	11.2	15.2	9.2
870_f_at	metallothionein 3 (growth inhibitory factor (neurotrophic))	MT3	M93311	90.7	15.1	42.9
56691_at			AI935202	26	15.1	1.2
39802_at	small inducible cytokine A7 (monocyte chemotactic protein 3)	SCYA7	X72308	1.8	15	10.2
875_g_at	small inducible cytokine A2 (monocyte chemotactic protein 1, homologous to mouse Sig-Je)	SCYA2	M26683	2.5	15	14.4
38369_at	myeloid differentiation primary response gene (88)	MYD88	U70451	5.1	15	18.4
45261_at	leucine aminopeptidase		AF034175	10.1	15	8.1

Figure 15J

34375_at	small inducible cytokine A2 (monocyte chemotactic protein 1, homologous to mouse Sig-je)	SCYA2	M28225	2.7	15	14.4
52696_at			AW024937	19.1	14.9	1.8
87258_at			N53015	4.1	14.9	9.8
48099_at			AA005023	13.7	14.8	4.6
42302_at			AI082042	55.4	14.7	1.9
70962_at			AA921754	0.3	14.7	0.5
48431_at	fucosyltransferase 4 (alpha (1,3) fucosyltransferase, myeloid-specific)	FUT4	AI970292	6	14.7	13.8
39108_at	lanosterol synthase (2,3-oxidosqualene-lanosterol cyclase)	LSS	U22526	17.1	14.6	4.4
51092_at	putative zinc finger protein NY-REN-34 antigen		AI950095	21.1	14.6	8.5
49163_at			N47894	2.2	14.6	4.7
57300_at	hypothetical protein FLJ11000		AI823982	15.6	14.6	3.5
35792_at	lysophospholipase-like		U67963	9.9	14.5	0
52905_at	Rag C protein		AI279898	26.9	14.5	19.7
38281_at	caspase 7, apoptosis-related cysteine protease	CASP7	U67319	22.7	14.4	4.9
384_at	proteasome (prosome, macropain) subunit, beta type, 10	PSMB10	X71874	7.2	14.4	1.2
71787_r_at			AI312358	19.4	14.4	0.5
64277_i_at	hypothetical protein FLJ20647		AI478759	7	14.4	9.8
53201_at			AI928513	61.2	14.4	1
39517_at	hypothetical protein 23851		AF035313	20.5	14.3	16.9
34745_at			AF070570	38.1	14.3	8.7
33513_at	signaling lymphocytic activation molecule	SLAM	U33017	178.3	14.3	1
56804_at			AA916789	12.3	14.3	3.6
69589_at			AI743005	15.4	14.3	42.7
32222_at	LR8 protein		AA152202	18.5	14.2	4
34474_at			W28429	1.1	14.2	0.4
61191_at			AI539443	3.9	14.2	9.4
49867_at			H44672	8.5	14.1	2
68047_at			AA936632	10.3	14	5.6
1729_at	TNFRSF1A-associated via death domain	TRADD	L41690	20.4	13.9	1.7
61873_at	glycerol kinase	GK	AI741715	20.7	13.9	0.5
60103_at			AI916948	46.9	13.9	44.2
43387_s_at	hypothetical protein FLJ13171		AW001604	6.4	13.8	9.9
53983_at	hypothetical protein PRO2714		AI139975	33.6	13.8	0.3

Figure 15K

38576_at	H2B histone family, member B	H2BFB	AJ223353	37.4	13.7	16.1
44673_at			N53555	9.5	13.7	4.3
55915_r_at	hypothetical protein LOC54149		AI922520	5.2	13.7	15.9
40670_at	kynureninase (L-kynurenine hydrolase)	KYNU	AI148772	52.5	13.6	7.9
40635_at	flotillin 1	FLOT1	AF089750	16.8	13.6	1.2
37360_at	lymphocyte antigen 6 complex, locus E	LY6E	U66711	13.4	13.6	8.3
51016_s_at	phosphoserine aminotransferase		AI971498	21.3	13.6	5.9
38160_at	lymphocyte antigen 75	LY75	AF011333	12.6	13.5	0
40687_at	gap junction protein, alpha 4, 37kD (connexin 37)	GJA4	M96789	74	13.5	4.2
52913_at			AA778521	11.5	13.5	0
73776_at			AI554044	18.6	13.5	14.8
55830_at			AA058686	19	13.5	8.8
36674_at	small inducible cytokine A4 (homologous to mouse Mip-1b)	SCYA4	J04130	30.3	13.4	13.3
46526_at	hypothetical protein MGC5242		AI434025	14.2	13.4	13.9
40700_at	nuclear body protein Sp140		U36500	3.7	13.3	19
34378_at	adipose differentiation-related protein	ADFP	X97324	5.1	13.3	0.1
36270_at	CD86 antigen (CD28 antigen ligand 2, B7-2 antigen)	CD86	U04343	4.9	13.2	3.9
34877_at	Janus kinase 1 (a protein tyrosine kinase)	JAK1	AL039831	18.4	13.2	2.9
51670_i_at			AI161358	45.6	13.2	52.1
31622_f_at	metallothionein 1F (functional)	MT1F	M10943	101.8	13.1	32.6
55673_at	RAB-8b protein		AI807023	16	13.1	6.6
45296_at	KIAA1631 protein		AI986035	12	13	4.3
53609_at			C14904	39.3	13	22.8
34749_at	solute carrier family 31 (copper transporters), member 2	SLC31A2	U83461	10.5	12.9	5.5
34699_at	CD2-associated protein	CD2AP	AL050105	20.4	12.9	8
47533_at			AI982765	18.6	12.9	14
33859_at	sin3-associated polypeptide, 18kD	SAP18	U96915	8	12.8	10.9
57501_at			AI914124	7.6	12.8	5.9
56080_at			AI703473	13.1	12.8	8.3
53855_at			N36284	20.1	12.7	12.5
61675_at			AI949662	9.1	12.7	14.2
80945_f_at			AI440145	15	12.7	2.5
64397_at	UCC1 protein		AA131797	2.1	12.7	1.6
52878_at			AI632502	0.3	12.7	30.9
41123_s_at	ectonucleotide pyrophosphatase/phosphodiesterase 2 (autotaxin)	ENPP2	L35594	1	12.6	1.8

Figure 15L

48080_at	hypothetical protein FLJ20037		AI566481	10.4	12.6	45.7
63066_at			AA001735	113	12.6	9.6
51988_at			AA131456	10.7	12.6	1.9
46152_at			AI760261	20.9	12.6	3
37348_s_at	thyroid hormone receptor interactor 7	TRIP7	AA845349	9.6	12.5	2.1
43269_at	KIAA1554 protein		AA424170	7.6	12.5	10.4
82071_at			AI138283	8.5	12.4	0
75026_s_at	methylene tetrahydrofolate dehydrogenase (NAD+ dependent), methenyltetrahydrofolate cyclohydrolase	MTHFD2	AI990317	6.9	12.4	0
45124_at			AI075999	10.8	12.4	0.8
36872_at	cyclic AMP phosphoprotein, 19 kD		AL120559	13.9	12.3	4.7
58317_at	KIAA1373 protein		AI638611	2.5	12.3	1.5
86261_at			AA017070	15.5	12.3	7.8
84995_at			AA769482	117.4	12.2	6.6
529_at	dual specificity phosphatase 5	DUSP5	U15932	45	12.1	10.5
36472_at	N-myc (and STAT) interactor	NMI	U32849	7.8	12.1	8.6
52167_at			AA151346	13.5	12.1	11.7
75000_at			AI735440	25.9	12.1	6.5
34304_s_at	spermidine/spermine N1-acetyltransferase	SAT	AL050290	17.7	12	9.3
64733_r_at	serine (or cysteine) proteinase inhibitor, clade B (ovalbumin), member 9	SERPINB9	AA156247	26	12	6.4
82933_at			AI312873	18.9	12	0.8
83734_at			AI381435	5.6	12	18.1
91306_s_at	ribosomal protein L37a	RPL37A	AA613715	105.7	12	0
79341_at			AA084622	7	12	17.6
37558_at	IGF-1I mRNA-binding protein 3		U97188	3.5	11.9	12.2
60017_at			AA142842	12	11.9	5.7
49339_s_at	reticulocalbin 1, EF-hand calcium binding domain	RCN1	AI817760	29.7	11.9	6.9
33352_at	H2A histone family, member Q	H2AFQ	X57985	8.1	11.8	37.7
38064_at	major vault protein	MVP	X79882	5	11.8	0.9
39230_at	similar to APOBEC1		AL022318	4.4	11.8	8.3
39168_at	Ac-like transposable element	ALTE	AB018328	1	11.8	23.5
63332_at			AA127696	19.8	11.7	11.4
79822_g_at			AI223310	1.6	11.7	2.4
52639_at			AA082625	3.3	11.7	4.8

Figure 15M

79721_at				AI733197	16.5	11.6	56.6
64344_at	lysosomal-associated membrane protein 1	LAMP1		AI984264	0.8	11.6	0.3
82588_at	CASP8 and FADD-like apoptosis regulator	CFLAR		AI302459	18.9	11.6	1.6
65540_at				AI863033	5.9	11.6	4.7
43839_f_at	glutaminase	GLS		AI982852	12.4	11.6	1.8
44640_at				W30985	40.2	11.6	8.4
45222_at	hypothetical protein FLJ12890			N23036	0	11.6	2.5
64938_at	hypothetical protein AL133206			AI702142	17.8	11.6	0.4
40082_at	fatty-acid-Coenzyme A ligase, long-chain 2	FACL2		D10040	16.9	11.5	6.5
408_at	GRO1 oncogene (melanoma growth stimulating activity, alpha)	GRO1		X54489	486.7	11.5	7.1
48768_s_at				R56038	8.8	11.5	1.1
595_at	tumor necrosis factor, alpha-induced protein 3	TNFAIP3		M59465	65.1	11.4	26.6
34300_at	zinc finger protein 278	ZNF278		AI352450	4.9	11.4	0.4
40362_at	nuclear factor of kappa light polypeptide gene enhancer in B-cells 2 (p49/p100)	NFKB2		X61498	29	11.4	5.2
50868_at				AW021051	1.2	11.4	7.2
50026_at	SAM domain, SH3 domain and nuclear localisation signals, 1	SAMSN1		AI823872	17.3	11.4	2.6
59516_at	KIAA1268 protein			AA210695	14.1	11.3	10.5
45234_at				AA648522	7.8	11.3	14.9
758_at	prostaglandin I2 (prostaglandin) receptor (IP)	PTGIR		D38128	24	11.3	2.6
52855_at				AI885781	4.7	11.3	14.6
46550_at	similar to mouse Xrn1 / Dhms2 protein			W01997	16	11.3	4.5
33260_at	son of sevenless (Drosophila) homolog 1	SOS1		L13857	12.8	11.2	5.3
48919_at	potassium inwardly-rectifying channel, subfamily J, member 2	KCNJ2		AA393850	130	11.2	13.1
36958_at	zyxin	ZYX		X95735	12.1	11.2	18.3
35149_at	tumor necrosis factor receptor superfamily, member 5	TNFRSF5		AI865431	15.8	11.1	6.1
85409_at				AI253672	10.9	11.1	12.7
44815_s_at				AA733119	92	11.1	28.6
58723_at				AA307796	10.4	11.1	5.3
609_f_at	metallothionein 1B (functional)	MT1B		M13485	116.3	11.1	20.1
56999_at				AI625959	28.4	11	3.4
75201_f_at				AI688900	5.5	11	7
37305_at	enhancer of zeste (Drosophila) homolog 2	EZH2		U61145	33.6	10.9	2.4
81887_at				AI766826	27.8	10.9	33.8
64333_s_at	pyrophosphatase (inorganic)	PP		AW009649	39.9	10.9	5

Figure 15N

72290_r_at				AA814901	9.1	10.9	12.8
49133_at	F-box only protein 6		FBXO6	AA167659	9.1	10.9	0.9
80482_at				AI247408	12	10.9	0.8
35153_at	Nijmegen breakage syndrome 1 (nibrin)		NBS1	AF058696	17.1	10.8	2.5
33924_at	KIAA1091 protein			AB029014	24.9	10.8	5.1
41516_at	SNARE protein			U95735	7.1	10.8	0.8
38342_at	KIAA0239 protein			D87076	4.2	10.8	16.9
38349_at				AF038564	12.9	10.8	0.5
50829_at				AA706818	4.8	10.8	10.5
74382_g_at				AA568587	24.4	10.8	0.8
54769_at				AA203416	33.9	10.8	14.7
65738_at	TERA protein			AA126787	19.3	10.8	1.4
77887_at				AL037345	0	10.8	0
1182_at	phospholipase C, epsilon		PLCE	D42108	0.3	10.7	8
35279_at	Tax1 (human T-cell leukemia virus type I) binding protein 1		TAX1BP1	U33821	9.3	10.7	0
36148_at	amyloid beta (A4) precursor-like protein 1		APLP1	U48437	14	10.7	2.1
41021_s_at	glycerol-3-phosphate dehydrogenase 2 (mitochondrial)		GPD2	U36310	16	10.7	9.1
81726_at				AI885498	15.1	10.7	7.2
53007_at				AA166622	8.7	10.7	1.5
36294_at	serine/threonine kinase 4		STK4	U60207	10	10.6	4.4
60019_at	leucine-rich neuronal protein		LRN	AA947309	13.9	10.6	1.4
62765_r_at	hypothetical protein FLJ22995			D20046	4.7	10.6	2.9
86629_at				AA614137	12.5	10.6	3.2
41503_at	KIAA0854 protein			AB020661	23.1	10.5	8.6
55763_at	MIL1 protein			AA225280	5.3	10.5	2.1
44729_at	hypothetical protein FLJ13855			AI027524	28.8	10.5	13.3
86122_l_at				AW015027	7.2	10.5	3.8
87651_at				AI800110	95.3	10.5	1
51104_at				AI937446	37.7	10.5	22.6
64393_at	KIAA1435 protein			AI695418	2	10.5	1.8
40932_at				H18080	3.8	10.4	1.7
33124_at	vaccinia related kinase 2		VRK2	AB000450	1.3	10.4	2.7
75868_r_at				AI791208	41.5	10.4	7
53720_at	hypothetical protein FLJ11286			AI862559	9.2	10.4	10.1
54720_at				AA493420	3.4	10.4	2

Figure 150

56805_at			N52761	16.8	10.4	14.1
53261_at			A1337231	34.9	10.4	1.6
1184_at	proteasome (prosome, macropain) activator subunit 2 (PA28 beta)	PSME2	D45248	9.3	10.3	5.7
906_at	signal transducer and activator of transcription 4	STAT4	L78440	54.6	10.3	5.9
56512_at			A1400772	9.6	10.3	0.2
82009_at			A1859620	10.6	10.3	1.6
50230_at			A1738719	12.4	10.3	3.2
38086_at	immunoglobulin superfamily, member 3	IGSF3	AB007935	1.4	10.2	3.3
44936_at			T93893	3.6	10.2	5.9
68328_at			A1475514	5.5	10.2	32.1
45204_s_at			A1378647	5	10.2	5.2
46612_at	KIAA1624 protein		R39794	5.2	10.2	1.6
41446_f_at	RNA helicase-related protein		H68340	71.4	10.1	21.9
49283_at			AA158597	19.1	10.1	0.9
54312_at			N34516	14.2	10.1	6
56641_at	liver-expressed antimicrobial peptide		A1937227	27	10.1	2.5
51510_r_at			T92882	18.1	10.1	4.2
37352_at	nuclear antigen Sp100	SP100	M60618	12.7	10	5
48929_at			AA521311	6.5	10	3.7
60839_at			A1924996	0.2	10	0
76098_r_at			A1304317	7.3	10	6
44853_s_at	CGI-34 protein		AA528059	12.9	10	7.4
62992_at			A1214996	25.6	10	4.3
40088_at	nuclear receptor interacting protein 1	NRIP1	X84373	15.6	9.9	8.5
848_at	TNF receptor-associated factor 1	TRAF1	U19261	33.5	9.9	5.5
32629_f_at	butyrophilin, subfamily 3, member A1	BTN3A1	U90552	10.8	9.9	8.4
36543_at	coagulation factor III (thromboplastin, tissue factor)	F3	J02931	32.3	9.9	4.2
1909_at	B-cell CLL/lymphoma 2	BCL2	M14745	12	9.9	8.1
59054_at			AA489100	4.1	9.9	4.8
42701_at			AA127763	9.4	9.9	0
64478_at			N36570	2.1	9.9	0.5
36671_at	asparagine synthetase	ASNS	M27396	16.8	9.8	3.2
36992_at	thioredoxin	TXN	A1653621	14.2	9.8	5.8
41048_at	phorbol-12-myristate-13-acetate-induced protein 1	PMAIP1	D90070	159.8	9.8	408.5
38352_at	peptidyl prolyl isomerase H (cyclophilin H)	PPIH	AF016371	3.2	9.8	1.2

Figure 15P

57772_g_at	hypothetical protein FLJ11200		AA886870	7.5	9.8	0
37420_i_at	major histocompatibility complex, class I, F	HLA-F	AL022723	12.3	9.8	2.6
57711_at			AA576959	78.3	9.8	0.7
80773_at			H15073	51.8	9.8	13.9
87135_at			AI239715	3.4	9.8	1.2
47483_at	similar to rat myomegalin		AI659612	71.1	9.8	11.8
39061_at	bone marrow stromal cell antigen 2	BST2	D28137	5.5	9.7	6.6
40556_at	reticulocalbin 1, EF-hand calcium binding domain	RCN1	D42073	14.7	9.7	5.4
41237_at	major histocompatibility complex, class I, A	HLA-A	D32129	14.9	9.7	5.6
1746_s_at	TNF receptor-associated factor 2	TRAF2	U12597	6.3	9.7	0
974_at	mitogen-activated protein kinase kinase 14	MAP3K14	Y10256	7	9.7	3.1
43939_at			AI590710	44.5	9.7	18
55629_at			AL040635	16.7	9.7	9.2
70228_r_at	protein kinase, cAMP-dependent, regulatory, type II, beta	PRKAR2B	AI792179	5.3	9.7	5
66025_at			AA075680	5.1	9.7	1.9
35095_r_at	leukocyte immunoglobulin-like receptor, subfamily A (without TM domain), member 3	LILRA3	AF025527	145.2	9.6	4.5
36149_at	dihydropyrimidinase-like 3	DPYSL3	D78014	2.3	9.6	1.2
80771_at			R73518	5.9	9.6	2.4
68716_at			AI674101	4	9.6	1.4
44561_r_at			AI096866	8.9	9.6	6.4
38357_at			AL049321	9.5	9.5	3
69800_at			AI806209	12.4	9.5	2.9
71617_at	hypothetical protein FLJ23112		AI937612	4	9.5	0
46188_at	hypoxia-inducible protein 2		AI422243	11.4	9.5	3.4
53970_at	dual adaptor of phosphotyrosine and 3-phosphoinositides		AF150266	43.8	9.5	3
89845_s_at	KIAA0084 protein		AI972597	11.3	9.5	0
75024_at	adenosine deaminase, RNA-specific	ADAR	R49062	7.6	9.5	4
38645_at	DKFZP434A043 protein		AL096748	1.1	9.4	2.1
34965_at	cystatin F (leukocystatin)	CST7	AF031824	17.7	9.4	2.7
41168_at	TAP binding protein (tapasin)	TAPBP	AF029750	9.2	9.4	2.4
39594_f_at	metallothionein 1H	MT1H	R93527	91.6	9.4	26.1
57051_at			AA167704	10.2	9.4	6.1
72228_at	chromodomain helicase DNA binding protein 2	CHD2	AI568867	12.4	9.4	0.2
47434_at			N24987	17.5	9.4	1.6

Figure 15Q

31812_at	guanosine monophosphate reductase		GMPR	M24470	1.1	9.3	5.5
37391_at	cathepsin L		CTSL	X12451	6.8	9.3	4.2
56373_at	UDP-Gal:betaGlcNAc beta 1,4- galactosyltransferase, polypeptide 5		B4GALT5	AA133969	42.2	9.3	9.5
51758_s_at				N21439	10	9.3	2.7
78732_f_at				AI094933	4.2	9.3	10.2
33484_at	ring finger protein 2		RNF2	Y10571	9.3	9.2	0
33305_at	serine (or cysteine) proteinase inhibitor, clade B (ovalbumin), member 1		SERPINB1	M93056	34.7	9.2	0
37723_at	cyclin G2		CCNG2	U47414	8.1	9.2	0
42345_at	KIAA1500 protein			N54300	3.1	9.2	4.3
65364_at				W56309	6.3	9.2	8.1
80332_at				AW006867	10.1	9.2	2.9
54585_at				AA927298	2.6	9.2	2.1
47596_at				AI066563	12	9.2	2.9
37137_at	granzyme B (granzyme 2, cytotoxic T-lymphocyte-associated serine esterase 1)		GZMB	M17016	52.3	9.1	15.3
37163_at	DKFZP586C1619 protein			AL050374	56.3	9.1	10.6
37853_at	urocortin		UCN	AI857458	21.7	9.1	9.9
33063_at	glycoprotein Ib (platelet), alpha polypeptide		GP1BA	J02940	30.9	9.1	9.4
38662_at				AL047596	6	9.1	4.5
37980_at	CBF1 interacting corepressor			U03644	4.1	9.1	8
46760_r_at				H38110	33.5	9.1	5.8
54975_at				AI761241	12.2	9.1	1.1
66951_at				AI016542	50.4	9.1	4.4
49302_at	HBV associated factor			AI971817	13.9	9.1	5.4
40468_at	KIAA0554 protein			AB011126	8.3	9	5.1
33852_at	TIA1 cytotoxic granule-associated RNA-binding protein		TIA1	M77142	6	9	1.9
38404_at	transglutaminase 2 (C polypeptide, protein-glutamine-gamma-glutamyltransferase)		TGM2	M55153	13.7	9	0.2
60835_at				AI831561	30.1	9	5.6
48531_at				AA522816	26	9	3.6
73176_r_at				AI249906	12	9	2.8
60016_at				AI400414	15.6	9	9.5
40054_at	KIAA0082 protein			D43949	8	8.9	1.8
1491_at	pentaxin-related gene, rapidly induced by IL-1 beta		PTX3	M31166	827.3	8.9	79

Figure 15R

36753_at	leukocyte immunoglobulin-like receptor, subfamily B (with TM and ITIM domains), member 4	LILRB4	AF072099	8.1	8.9	3.9
76762_at			AI246590	53.8	8.9	10.2
66084_at			T81819	0	8.9	5.5
46296_at	hypothetical protein FLJ10579		W60903	2.4	8.9	4.9
37047_at	Niemann-Pick disease, type C1	NPC1	AF002020	0.9	8.8	1.4
79520_at			AW022213	18.4	8.8	8
86536_at			AI740796	16.4	8.8	1.7
54933_at			AI356461	11.3	8.8	4.7
57674_at	Fanconi anemia, complementation group F	FANCF	AI978840	5.7	8.8	3.1
43848_s_at			AI968465	9.1	8.8	27.3
57697_at			AI445628	0	8.8	0.8
48097_at	transmembrane protein with EGF-like and two follistatin-like domains 1	TMEFF1	AA001144	13.2	8.8	0
46317_at	CD36 antigen (collagen type I receptor, thrombospondin receptor)-like 2 (lysosomal integral membrane protein II)	CD36L2	W07476	0.8	8.8	10.5
38149_at	KIAA0053 gene product		D29642	9.1	8.7	4.3
1617_at	zinc finger protein 147 (estrogen-responsive finger protein)	ZNF147	D21205	1.8	8.7	0.7
195_s_at	caspase 4, apoptosis-related cysteine protease	CASP4	U28014	11.6	8.7	4.5
36185_at	alanyl-tRNA synthetase	AARS	D32050	14.4	8.7	0
37218_at	BTG family, member 3	BTG3	D64110	9.7	8.7	6.5
37544_at	nuclear factor, interleukin 3 regulated	NFIL3	X64318	8.9	8.7	3.2
43901_i_at			AI435160	20.5	8.7	2
63664_at			AA186367	12.5	8.7	2.3
73235_at	guanylate binding protein 1, interferon-inducible, 67kD	GBP1	AW014593	13.7	8.7	1.7
54565_at			AA149736	45.6	8.7	10.3
317_at	protease, cysteine, 1 (legumain)	PRSC1	D55696	2.5	8.6	7.2
1358_s_at	interferon, alpha-inducible protein (clone IFI-6.16)	G1P3	U22970	11.3	8.6	10
33328_at			W28612	1.7	8.6	11.7
65139_at	hypothetical protein FLJ12150		AI925240	4.5	8.6	0.8
84981_r_at	collagen, type III, alpha 1 (Ehlers-Danlos syndrome type IV, autosomal dominant)	COL3A1	T54146	5.1	8.6	5.6
81932_at			AI659966	16.5	8.6	0
65774_at			AI556963	0.4	8.6	0
64174_at			AI127460	11	8.6	7

Figure 15S

56871_at				AI953847	25.7	8.6	12.8
45785_at	DKFZP566B1346 protein			AL040982	19.7	8.6	3.4
63982_at	ubiquinol-cytochrome c reductase (6.4kD) subunit			AI480328	29.8	8.6	9.8
32065_at	cAMP responsive element modulator	CREM		S68134	7.7	8.5	47.4
34777_at	adrenomedullin	ADM		D14874	27.2	8.5	7.1
39988_at	ATP-binding cassette, sub-family B (MDR/TAP), member 3	ABCB3		M74447	15.9	8.5	3
39993_at	phosphatidylinositol glycan, class A (paroxysmal nocturnal hemoglobinuria)	PIGA		D11466	16.9	8.5	5.8
35337_at	F-box only protein 7	FBXO7		AL050254	8.6	8.5	16.3
1665_s_at	endothelial cell growth factor 1 (platelet-derived)	ECGF1		M63193	7.3	8.5	1.6
37616_at	AU RNA-binding protein/enoyl-Coenzyme A hydratase	AUH		X79888	7	8.5	0
38315_at	retinaldehyde dehydrogenase 2			AB015228	17.3	8.5	0.4
57537_at				AA001250	9.6	8.5	4.6
60288_at				AA465415	1.9	8.5	1.6
58288_at				W63676	2.3	8.5	1.9
81650_at				AA631047	1.9	8.5	0
69266_at				AI634394	1.7	8.5	3.6
57184_at				AI268299	2.8	8.5	0
53489_at				AI928764	18.3	8.5	20.8
46858_at				AI796221	8.5	8.5	4.4
52337_g_at				AI762208	28	8.5	7.5
57733_at				AI652899	18.5	8.5	1.6
319_g_at	H1 histone family, member X	H1FX		D64142	12.9	8.4	0.7
40522_at	glutamate-ammonia ligase (glutamine synthase)	GLUL		X59834	7	8.4	0
1446_at	proteasome (prosome, macropain) subunit, alpha type, 2	PSMA2		D00760	8.9	8.4	2.6
38465_at	peptidylglycine alpha-amidating monooxygenase	PAM		M37721	11.9	8.4	0.1
40698_at	C-type (calcium dependent, carbohydrate-recognition domain) lectin, superfamily member 2 (activation-induced)	CLECSF2		X96719	0.3	8.4	6.5
50855_at				AA789332	0	8.4	7
52721_r_at				AA618532	6	8.4	5.7
70594_at				AI819198	102.7	8.4	2.5
55843_at	Jun dimerization protein p21SNFT			AI359117	21	8.4	0.1
57266_r_at	pleiomorphic adenoma gene-like 1	PLAGL1		AI971679	9.2	8.4	1.8
60424_at	hypothetical protein DKFZp761B2423			AA971604	28.4	8.4	58
47572_at				AI927902	7.6	8.4	0.9

Figure 15T

41409_at	basement membrane-induced gene		AF044896	12.7	8.3	0
35879_at	galanin	GAL	M77140	5.4	8.3	0.4
40936_at	cysteine-rich motor neuron 1	CRIM1	AI651806	9.6	8.3	0.9
33849_at	pre-B-cell colony-enhancing factor		U02020	41	8.3	16.3
41164_at	immunoglobulin heavy constant mu	IGHM	X67301	8.9	8.3	47.2
35302_at	nuclear RNA export factor 1	NXF1	AJ132712	6.9	8.3	34.6
37536_at	CD83 antigen (activated B lymphocytes, immunoglobulin superfamily)	CD83	Z11697	10.7	8.3	6.3
61118_at			AI983632	13.4	8.3	3.1
74482_at			AW025586	1.2	8.3	7.4
60968_at			AI990212	25.3	8.3	4.9
46961_r_at			T91186	6.9	8.3	9.6
37126_at	Sjogren syndrome antigen A1 (52kD, ribonucleoprotein autoantigen SS-A/Ro)	SSA1	M62800	8.6	8.2	10.6
34678_at	fer-1 (C.elegans)-like 3 (myoferlin)	FER1L3	AL096713	3.3	8.2	3.4
37312_at	KIAA0127 gene product		D50917	7.8	8.2	1.2
43581_at			AA262338	3.1	8.2	0
73455_f_at			H72723	70.4	8.2	8.5
56697_r_at			AI791520	7.1	8.2	8.3
40570_at	forkhead box O1A (rhabdomyosarcoma)	FOXO1A	AF032885	6.1	8.1	4.5
377_g_at	sema domain, immunoglobulin domain (lg), short basic domain, secreted, (semaphorin) 3C	SEMA3C	AB000220	5.5	8.1	0
33736_at	stomatin (EBP72)-like 1	STOML1	Y16522	8.8	8.1	1.8
41329_at	hypothetical protein LOC57147		AI458463	7.6	8.1	5.7
404_at	interleukin 4 receptor	IL4R	X52425	9	8.1	1.5
56458_at			Z99408	0	8.1	0.8
90732_at	splicing factor, arginine/serine-rich 5	SFRS5	W45137	2.1	8.1	3
43038_r_at	KIAA1363 protein		AI769531	0	8.1	4
62101_g_at	anaphase-promoting complex 1, meiotic checkpoint regulator		N90510	9.3	8.1	0.7
60188_at			AA629050	4.5	8.1	3.1
46120_at			AA149250	2	8.1	0
77840_f_at			AI720888	4.4	8.1	16.6
32218_at			AF034176	8.1	8.1	5.5
40451_at	polymerase (DNA directed), epsilon	POLE	AL080203	11.2	8	0
36493_at	lymphocyte-specific protein 1	LSP1	M33552	0	8	0

Figure 15U

36573_at	DEAD/H (Asp-Glu-Ala-Asp/His) box binding protein 1	DDXBP1	U78524	3.7	8	6.6
40755_at	MHC class I polypeptide-related sequence A	MICA	X92841	15.4	8	13.9
54744_at			AA582287	23.5	8	3
75183_at	signal transducer and activator of transcription 5A	STAT5A	AI990633	15.3	8	0
54857_at	ancient conserved domain protein 4		AI636759	9.6	8	7.1
86863_at			AI798714	10.1	8	0.5
46175_i_at			AA552969	22	8	3
46655_at	transcription factor (SMIF gene)		AA131327	16.6	8	14.5
56769_at	B7-H1 protein		AA292201	25.6	8	6.1
32128_at	small inducible cytokine subfamily A (Cys-Cys), member 18, pulmonary and activation-regulated	SCYA18	Y13710	7.8	7.9	4
41126_at	solute carrier family 1 (glutamate/neutral amino acid transporter), member 4	SLC1A4	AA978353	5.7	7.9	4.7
34887_at			N92548	14.3	7.9	1.4
57663_at			AA600810	1.8	7.9	1.4
62431_at			AI197909	7.1	7.9	7.6
67382_at			AA936421	4	7.9	4
52273_s_at	RAB10, member RAS oncogene family	RAB10	AI979258	3.6	7.9	2.9
53578_at			W70067	17.9	7.9	3.9
78264_at			N49445	1.7	7.9	1.4
64484_at			AW006123	5.8	7.9	5
48932_r_at			AA059347	1.3	7.9	3.3
48753_at	PPAR(gamma) angiotensin related protein		N57259	126	7.9	14.7
39861_at	ems1 sequence (mammary tumor and squamous cell carcinoma-associated (p80/85 src substrate)	EMS1	M98343	8.4	7.8	6.2
894_g_at	ubiquitin carrier protein		M91670	5.9	7.8	5.4
38555_at	dual specificity phosphatase 10	DUSP10	AB026436	18.8	7.8	11.2
34256_at	sialyltransferase 9 (CMP-NeuAc:lactosylceramide alpha-2,3-sialyltransferase; GM3 synthase)	SIAT9	AB018356	0	7.8	5.7
39660_at	defensin, beta 1	DEFB1	AI309115	21.1	7.8	15.1
31540_at	tumor necrosis factor receptor superfamily, member 9	TNFRSF9	U03397	72.2	7.8	19.5
55257_at	prion gene complex, downstream		W76645	1.3	7.8	5.8
53727_at			AI768608	20	7.8	1.7
90083_at			AA149818	11.2	7.8	0
81035_i_at			AI573275	14.8	7.8	1.9

Figure 15V

67223_at				AI003757	6.9	7.8	6.5
83874_at				AI024818	23.9	7.8	4.4
69142_at				AI635522	72.4	7.8	8.6
46104_at				AA772055	0.2	7.8	6.7
45838_at				H12244	6.4	7.8	9.2
47055_at				AA481246	1.1	7.8	0
39211_at				AF091080	12.5	7.7	5.7
37324_at			TFRC	X01060	13	7.7	1.3
38091_at			LGALS9	Z49107	3.2	7.7	6.7
38450_at			SSB	X69804	1.4	7.7	16.2
56534_at				N20945	1.8	7.7	0
85561_at				AL079435	76.5	7.7	0.6
63013_at				AL120446	6.2	7.7	4.2
48773_at				AA642981	10.3	7.7	0
38970_s.at				AJ011896	26.4	7.6	4.2
34530_at				W73822	4.5	7.6	1.2
40452_at			CPNE1	U83246	9.5	7.6	0.9
33905_at			MBD2	AF072242	6.1	7.6	0.5
36564_at				W27419	24.2	7.6	2
37363_at				AB007889	7.9	7.6	3.8
37625_at			IRF4	U52682	6.5	7.6	1.7
570_at			RELB	M83221	11.9	7.6	4
82055_at				AL119129	9.5	7.6	1.6
79371_at				AA203372	9.4	7.6	3
43829_at				AW003197	12.4	7.6	0.3
50693_at				2836883	6.5	7.6	1.1
41678_at			EPHB2	AF025304	0.8	7.5	4.7
1383_at			PPP2R2A	M64929	5.3	7.5	12.4
36526_at			EXTL2	AF000416	22.5	7.5	0
38372_at				U66042	3	7.5	0
38439_at			NFE2L1	L24123	12	7.5	3.5
77058_at				AA044778	4.1	7.5	2.8
78538_at				AI023295	5.4	7.5	2

Figure 15W

46697_at	protein kinase, AMP-activated, gamma 2 non-catalytic subunit	PRKAG2	AA115330	1.3	7.5	6.9
48076_at	hypothetical protein		AI949434	24.1	7.5	19.9
56039_at			AI718675	1.4	7.5	2.4
44978_at	Ac-like transposable element	ALTE	AI979194	1	7.5	12.7
31850_at	glutamate-cysteine ligase, catalytic subunit	GCLC	M90656	1.3	7.4	32.4
39628_at	RAB9, member RAS oncogene family	RAB9	AI671547	12.8	7.4	3.5
32069_at	KIAA0615 gene product		AB014515	13.5	7.4	10
33977_at	growth factor independent 1	GFI1	U67369	0.9	7.4	0.9
38014_at	adenosine deaminase, RNA-specific	ADAR	X79448	7.8	7.4	5.9
49136_at			W56900	1.1	7.4	12.3
59061_at			AI670069	9.4	7.4	8.4
55239_at			AA233120	6.1	7.4	6.8
85241_at			AA846371	1	7.4	0
49496_at			AA521168	1.1	7.4	17.1
52675_at			AI581142	16.5	7.4	3.6
49650_at			N68715	9.8	7.4	3.1
53139_at	protein phosphatase 1B (formerly 2C), magnesium-dependent, beta isoform	PPM1B	AW020971	10.6	7.4	11.9
46034_r_at			AA074612	7.2	7.4	2.6
48742_at	activated RNA polymerase II transcription cofactor 4		AI979012	7.3	7.4	3.6
34214_at	KIAA0644 gene product		AB014544	2.5	7.3	20.7
1035_g_at	tissue inhibitor of metalloproteinase 3 (Sorsby fundus dystrophy, pseudoinflammatory)	TIMP3	U14394	1.5	7.3	0.6
33524_at	v-crk avian sarcoma virus CT10 oncogene homolog-like	CRKL	X59656	1	7.3	0.4
35321_at	tousled-like kinase 2	TLK2	AB004884	7.8	7.3	4.5
32863_at	similar to calcium/calmodulin dependent protein kinases		AL023754	28.7	7.3	1.9
48021_at	kinesin heavy chain member 2	KIF2	AI652174	6.2	7.3	2.8
62512_at			AI927692	9.7	7.3	18.4
53141_at			AI810396	8.1	7.3	0
48081_at	myristoylated alanine-rich protein kinase C substrate (MARCKS, 80K-L)	MACS	AA131320	18.7	7.3	8.2
89856_at	CD83 antigen (activated B lymphocytes, immunoglobulin superfamily)	CD83	AI983994	7.3	7.3	5.3
44884_at			AA039908	4.1	7.3	6.8
40619_at	ubiquitin carrier protein		M91670	2.8	7.2	3.8

Figure 15X

32224_at	KIAA0769 gene product			AB018312	5.9	7.2	0.5
33759_at	2,3-bisphosphoglycerate mutase	BPGM		X04327	2	7.2	1
33418_at	RAB3 GTPase-ACTIVATING PROTEIN			AL096752	5.6	7.2	5.2
34759_at				U68494	3.9	7.2	23.3
35764_at	oral-facial-digital syndrome 1 gene	OFD1		Y15164	2	7.2	2.4
36833_at	Bruton agammaglobulinemia tyrosine kinase	BTK		U78027	4.1	7.2	1.1
1402_at	v.yes.1 Yamaguchi sarcoma viral related oncogene homolog	LYN		M16038	10.4	7.2	5.2
38472_at	KIAA0143 protein			D63477	0.3	7.2	3.1
40712_at	a disintegrin and metalloproteinase domain 8	ADAM8		D26579	4.9	7.2	1.2
62217_at				T81422	23.7	7.2	15.8
89178_at				AI741934	0	7.2	6.5
50427_at				AW003857	10.5	7.2	3.5
45390_at				AI769269	5.5	7.2	0
45841_at				AI916641	0	7.2	12
32805_at	aldo-keto reductase family 1, member C1 (dihydrodiol dehydrogenase 1; 20-alpha (3-alpha)-hydroxysteroid dehydrogenase)	AKR1C1		U05861	8.8	7.2	0
44696_at	hypothetical protein FLJ10743			AA915989	7.1	7.2	0
64103_at				N22751	3.1	7.2	9.1
64119_at				AA195074	2.5	7.2	0.2
73128_at	dihydropyrimidinase-like 3	DPYSL3		AI983251	5	7.2	0
75246_s_at	solute carrier family 1 (glial high affinity glutamate transporter), member 3	SLC1A3		W26838	43.1	7.2	4.8
51508_i_at				T92882	13.7	7.2	9.7
39541_at	KIAA1237 protein			W52003	2.7	7.1	9.7
39821_s_at	growth arrest and DNA-damage-inducible, beta	GADD45B		N95168	11.8	7.1	21.4
40215_at	UDP-glucose ceramide glucosyltransferase	UGCG		D50840	6	7.1	1.7
35166_at	Down syndrome critical region gene 3	DSCR3		D87343	4.6	7.1	5.7
77680_at				AI809979	3.9	7.1	3.1
47500_i_at	c-myc promoter-binding protein			AA805337	20.6	7.1	2.8
84765_at				AI446030	4.4	7.1	1.9
55551_at				AI766561	10.6	7.1	10.3
45256_at	pre-B-cell colony-enhancing factor			AI271460	35.1	7.1	33.4
54321_at				AI290473	9.7	7.1	2.3
51813_at				AI797095	4.8	7.1	1.8

Figure 15Y

56224_at	chloride channel, nucleotide-sensitive, 1A	CLNS1A	N32865	7.6	7.1	0
56866_at			AA565426	4.2	7.1	1.9
69573_at			A1718148	16.9	7.1	3.6
38649_at	KIAA0970 protein		AB023187	14.2	7	30.1
1310_at	proteasome (prosome, macropain) subunit, beta type, 2	PSMB2	D26599	5.5	7	4.6
41776_at	ATX1 (antioxidant protein 1, yeast) homolog 1	ATOX1	U70660	1.6	7	0.6
40290_f at	sialyltransferase 4A (beta-galactosidase alpha-2,3-sialyltransferase)	SIAT4A	L13972	3	7	2.2
35704_at	similar to rat HREV107		X92814	2.5	7	0.5
36175_s at	human immunodeficiency virus type I enhancer-binding protein 2	HIVEP2	AL023584	24.2	7	5.7
37716_at	antigen identified by monoclonal antibody MRC OX-2	MOX2	X05323	50.5	7	0
63683_at			N56920	7	7	11
58944_s at	ribosomal protein S6 kinase, 52kD, polypeptide 1	RPS6KC1	AA670286	30	7	5.6
53498_at			AA464480	9.6	7	6.1
53665_at			A1830209	9.2	7	1.6
74680_r at	FLN29 gene product		AW002527	4.7	7	6.3
68855_r at			AA594352	6.5	7	3.9
37485_at	fatty-acid-Coenzyme A ligase, very long-chain 1	FACVL1	D88308	1.5	6.9	6.5
269_at	2'-5'-oligoadenylate synthetase-like	OASL	L40387	19.8	6.9	119.4
289_at	signal transducer and activator of transcription 3 (acute-phase response factor)	STAT3	L29277	13.6	6.9	2.2
1170_at	colony stimulating factor 1 (macrophage)	CSF1	M37435	13.1	6.9	3.2
33253_at	KIAA0129 gene product		D50919	0	6.9	0.1
39320_at	caspase 1, apoptosis-related cysteine protease (interleukin 1, beta, convertase)	CASP1	U13697	5.1	6.9	1.1
32233_at	torsin family 1, member B (torsin B)	TOR1B	AF007872	5	6.9	13
33285_i at			W26762	3.3	6.9	0
41520_at	hypothetical protein		AC005757	10.2	6.9	2.9
36456_at	DKFZP564I052 protein		AL080063	10.7	6.9	4.7
35340_at	mel transforming oncogene (derived from cell line NK14): RAB8 homolog	MEL	A1819948	0	6.9	10.5
37971_at	bromodomain adjacent to zinc finger domain, 1A	BAZ1A	AL050089	6.7	6.9	6.4
56832_at			A1871408	13.7	6.9	12.3
80193_at			A1130878	7.1	6.9	0
59365_at			D59637	32.9	6.9	4.9
63398_at	similar to Bos taurus P14 protein		AA130220	4.4	6.9	2.1

Figure 15Z

58825_at	prothymosin, alpha (gene sequence 28)	PTMA	AI701911	5.4	6.9	7.3
64176_at			AI650542	10.3	6.9	8.5
42365_f_at			H60126	10.8	6.9	3.3
45624_at	core1 UDP-galactose:N-acetylgalactosamine-alpha-R beta 1,3-galactosyltransferase		AI967968	5.2	6.9	4
55353_at			AA760977	33.6	6.9	6
44120_at			AI879381	6.9	6.9	7.4
40218_at	CDP-diacylglycerol synthase (phosphatidate cytidyltransferase) 1	CDS1	U60808	0.1	6.8	6.7
32331_at	adenylate kinase 3	AK3	X60673	42.3	6.8	1
90103_at			AA503824	7.6	6.8	4
77119_at			AI820859	10	6.8	1.2
58774_at	eukaryotic translation initiation factor 4 gamma, 1	EIF4G1	AI380979	0.7	6.8	0
60271_at	hypothetical protein similar to small G proteins, especially RAP-2A		AA830857	31.8	6.8	2.5
44055_at			AA133979	0	6.8	9.6
65554_at			AI188518	3.4	6.8	3.3
88103_s_at	hypothetical protein FLJ20608		AW007018	6.1	6.8	0.1
38459_g_at	cytochrome b-5	CYB5	L39945	3.4	6.8	5.6
48327_at			AA831325	2.6	6.8	4
91719_at			AA974392	1.3	6.8	2.4
37684_at	solute carrier family 21 (organic anion transporter), member 9	SLC21A9	AB020687	8.4	6.8	8
33380_at	topoisomerase-related function protein 4-1		AB005754	6.9	6.7	20
36052_at	adducin 2 (beta)	ADD2	U43959	1.1	6.7	4.4
32250_at	H factor 1 (complement)	HF1	X07523	2.6	6.7	4.6
36770_at	signal transducer and activator of transcription 2, 113kD	STAT2	U18671	4.5	6.7	6.7
37046_at	proteasome (prosome, macropain) subunit, alpha type, 5	PSMA5	AI246726	9.8	6.7	2.6
39138_g_at	nuclear factor related to kappa B binding protein	NFRKB	X80878	0.7	6.7	3.2
58005_at	hypothetical protein		AI480253	3.7	6.7	3.8
67295_at			AI362842	9.5	6.7	2.3
49345_at			AI770110	1.1	6.7	3.3
42931_at			W87642	6.1	6.7	5.8
64047_at	KIAA1376 protein		AA587245	26.1	6.7	38.4
75525_at	KIAA0336 gene product		R34331	4.6	6.7	2.1
52479_r_at			AI888055	2.9	6.7	1.2
46260_at	claudin 1	CLDN1	AI452474	9.2	6.7	2.4
63027_at			AA151719	3.1	6.7	3.4

Figure 15AA

88499_f at				AI650658	0.6	6.7	0.8
39702_at	ribonuclease P (30kD)			U77665	6.6	6.7	6.2
31492_at	muscle specific gene			AB019392	8.5	6.7	9.5
44688_at	RelA-associated inhibitor			AA614049	7.7	6.7	8.8
41670_at	hypothetical protein			R38263	8.1	6.6	0.4
2009_at	protein tyrosine kinase 2 beta		PTK2B	U33284	4.8	6.6	4.3
32158_at	RAD9 (S. pombe) homolog		RAD9	U53174	2.8	6.6	5.5
41343_at	CDP-diacylglycerol synthase (phosphatidate cytidylyltransferase) 2		CDS2	Y16521	9.5	6.6	0
40612_at	KIAA1117 protein			AB029040	4.6	6.6	3.6
514_at	Cas-Br-M (murine) ectropic retroviral transforming sequence b		CBLB	U26710	16.4	6.6	1.5
1860_at	tumor protein p53-binding protein, 2		TP53BP2	U58334	4.6	6.6	0.5
566_at	promyelocytic leukemia		PML	M79462	6.1	6.6	8.4
39158_at	activating transcription factor 5		ATF5	AB021663	6.7	6.6	4.7
90388_at				AA909008	1.5	6.6	7.9
82402_at	hypothetical protein DKFZp761D1823			AA522440	0	6.6	0
75645_at				AI795971	1.1	6.6	2.7
67711_s_at				AI986192	9.6	6.6	0.1
87957_r_at				AA513538	13.9	6.6	3.6
49516_at				AA531016	5	6.6	3.5
86168_at				AA745606	2.2	6.6	33.8
46246_at	N-Acetylglucosamine kinase			AW026312	0	6.6	0.7
48276_at				AI275691	19.9	6.6	2
747_at				D63134	6.5	6.6	1.2
34926_at	CD1A antigen, a polypeptide		CD1A	M28825	7.1	6.6	5.7
824_at	glutathione-S-transferase like; glutathione transferase omega			U90313	4.6	6.5	0
39723_at	cullin 1		CUL1	AF062536	16	6.5	4.8
40458_at	signal transducer and activator of transcription 5A		STAT5A	U43185	8.8	6.5	1.9
40907_at				AA043502	6.4	6.5	8.3
33741_at	CGI-11 protein			AI741756	0.8	6.5	3
37177_at	CD58 antigen, (lymphocyte function-associated antigen 3)		CD58	Y00636	5.6	6.5	0.6
48105_at	latexin protein			AI672094	27.4	6.5	0.9
56428_at	hematological and neurological expressed 1		HN1	AI525822	1	6.5	0.7
49961_at	KIAA0477 gene product			AI821791	11.3	6.5	0
80847_at				AA352113	4.4	6.5	1.4
80896_at				AA602921	5.8	6.5	2.1

Figure 15BB

38567_at	CD1D antigen, d polypeptide	CD1D	L38820	2.4	6.5	0.5
50761_at	BANP homolog, SMAR1 homolog		AI738764	3.8	6.5	1.1
50266_at			AI741178	7.7	6.5	5.4
75247_f_at			AA548037	3.4	6.5	4.6
47655_at			AI022328	6.4	6.5	8.6
44149_at			AI674647	8.4	6.5	1.5
55361_at	thromboxane A synthase 1 (platelet, cytochrome P450, subfamily V)		AI821565	4.3	6.5	8.3
33777_at	RAB13, member RAS oncogene family	TBXAS1	D34625	6.6	6.5	6.3
40210_at	basic leucine zipper nuclear factor 1 (JEM:1)	RAB13	X75593	14.1	6.4	0.6
32088_at		BLZF1	U79751	9.7	6.4	16.3
38207_at	KIAA0471 gene product		AW006742	13.4	6.4	12.4
34445_at	chemokine (C-C motif) receptor-like 2		AB007940	1.2	6.4	16.7
1445_at	human immunodeficiency virus type 1 enhancer-binding protein 1	CCR12	AF014958	32.3	6.4	3.5
35251_at		HIVEP1	X51435	5.6	6.4	12.2
36282_at	protein tyrosine phosphatase, non-receptor type 2	PTPN2	AF054994	1.4	6.4	9
1265_g_at			M25393	24.4	6.4	1.2
42524_at			R37337	30.1	6.4	3.6
82094_i_at			AI810266	12.4	6.4	36.3
65329_at			H56582	0	6.4	1.8
48556_at	HIV TAT specific factor 1	HTATSF1	AA581365	16.4	6.4	4.2
56965_at	HSPC056 protein		N51105	4.9	6.4	1
58933_i_at	hypothetical protein FLJ20446		AA526392	0.7	6.4	14.2
48566_at	leucine-rich protein mRNA		N98615	7.7	6.4	3.8
70516_at	mitogen-activated protein kinase kinase kinase 13		AI817736	14.2	6.4	8.6
49450_at	hypothetical protein FLJ10895	MAP3K13	W72274	11	6.4	0
46371_at	dynactin 1 (p150, Glued (Drosophila) homolog)		N36090	1.2	6.4	6.4
51865_at		DCTN1	W80446	1.5	6.4	2.7
91390_at	KIAA0430 gene product		AA508196	8.1	6.4	0
64508_s_at	STAT induced STAT inhibitor-2		AI968969	7.4	6.4	8.5
42141_at	interferon-stimulated transcription factor 3, gamma (48kD)		AA469994	3	6.3	2
38994_at		ISGF3G	AF037989	29.2	6.3	1.8
38517_at	basic leucine zipper transcription factor, ATF-like		M87503	10	6.3	11.4
32217_at	KIAA0084 protein	BATF	AF052105	12.6	6.3	7.6
39942_at			AF016898	25.6	6.3	0
32593_at			D42043	12.5	6.3	0.5

Figure 15CC

48517_at	GTP binding protein 2	GTPBP2	H88399	7.9	6.3	0
91895_f_at			A1335251	4.1	6.3	3.2
344_s_at	2',3'-cyclic nucleotide 3' phosphodiesterase	CNP	D13146	15.9	6.3	1.6
80964_at			AA927180	36.3	6.3	0
91984_r_at			A1479022	3	6.3	2.3
70151_at			A1805522	9.8	6.3	5.3
56278_at			A1692445	15.4	6.3	58
65764_at	RAB18, member RAS oncogene family	RAB18	D12273	3.4	6.3	0.3
71643_at			A1917799	1.3	6.3	1.7
66092_at			A1951574	6.2	6.3	6.5
43553_at			A1807404	7.2	6.3	3.1
60857_at	chloride intracellular channel 4	CLIC4	AA142983	20.5	6.3	2.7
86670_at			AA604317	11.3	6.3	0.5
47971_at	ferritin, heavy polypeptide 1	FTH1	T92243	27.6	6.3	0.8
51854_at	KIAA1128 protein		AA424126	14.4	6.3	5.6
48732_at			W73033	7.3	6.3	3
51028_at			A1218306	2.6	6.3	4.7
65827_at	STIP1 homology and U-Box containing protein 1	STUB1	A1630895	7.2	6.3	8.1
39210_at	fucosyltransferase 4 (alpha (1,3) fucosyltransferase, myeloid-specific)	FUT4	M58597	2.8	6.2	8.8
1449_at	proteasome (prosome, macropain) subunit, alpha type, 4	PSMA4	D00763	8.3	6.2	10.1
1953_at	vascular endothelial growth factor	VEGF	AF024710	28.4	6.2	7.2
91669_at			AA491645	8.4	6.2	5.6
82421_i_at			A1554417	4.1	6.2	8.8
59451_at			A1800640	5.3	6.2	23.2
44048_at	karyopherin (importin) beta 2	KPNB2	AA527296	3.4	6.2	0
56246_at	6-phosphofructo-2-kinase/fructose-2,6-biphosphatase 3	PFKFB3	A1765775	91	6.2	11.3
50388_at	MFH-amplified sequences with leucine-rich tandem repeats 1		AA044708	9.8	6.2	2.2
45491_s_at	myeloid/lymphoid or mixed-lineage leukemia (trithorax (Drosophila) homolog); translocated to, 6	MLLT6	A1828593	6.5	6.2	1.9
51155_at			AL043669	19.8	6.2	8.7
52186_at	niban protein		A1560064	11.7	6.2	1.7
46153_at			R02594	5.7	6.2	4
46183_at			A1810826	7	6.2	4.6
55991_at			AA521262	10.9	6.2	1.4

Figure 15DD

58682_at	DEAD/H (Asp-Glu-Ala-Asp/His) box binding protein 1	DDXBP1	AI348378	4.8	6.2	0
54982_at	pellino (Drosophila) homolog 1	PELI1	AL043980	85.4	6.2	54.5
57543_at	bromodomain adjacent to zinc finger domain, 1A	BAZ1A	AA112166	8.3	6.2	6.5
38712_at	membrane protein CH1		AL035291	6.7	6.1	6.9
41635_at	Wilms' tumour 1-associating protein		D14661	31.6	6.1	18.4
31859_at	matrix metalloproteinase 9 (gelatinase B, 92kD gelatinase, collagenase)	MMP9	J05070	1	6.1	0.4
39221_at	leukocyte immunoglobulin-like receptor, subfamily B (with TM and ITIM domains), member 2	LILRB2	AF004231	62.6	6.1	17.8
36103_at	small inducible cytokine A3 (homologous to mouse Mip-1a)	SCYA3	D90144	13.1	6.1	7.6
37673_at	neutral sphingomyelinase (N-SMase) activation associated factor	NSMAF	X96586	6.2	6.1	0
37679_at	interferon-related developmental regulator 1	IFRD1	Y10313	20.1	6.1	2.1
90428_at			AI355272	6.2	6.1	3.4
82077_at			AI916280	7.7	6.1	3.5
62266_at	hypothetical protein FLJ21079		AI339505	14.7	6.1	1.7
82594_at			AI308862	5.8	6.1	3.1
48528_at			Z99386	10.2	6.1	0
54819_r_at			AI720898	16.6	6.1	1.3
64007_at	KIAA0964 protein		AI471030	9	6.1	0
43810_at	KIAA0769 gene product		AW003036	7.4	6.1	0
69197_r_at			AA508552	21.6	6.1	0
89210_at			AI741905	12	6.1	2.7
72808_at			AI640523	16.1	6.1	32.6
44069_at			H03287	2.7	6.1	3.5
43592_at	chondroitin 4-O-sulfotransferase 2		AA449104	0.5	6.1	10.6
48822_s_at	adenylate kinase 3	AK3	AW015546	41.5	6.1	0.5
44677_at	AD-015 protein		N42213	8.3	6.1	0
51030_at			AA843962	13.3	6.1	2.2
47321_at	a disintegrin-like and metalloprotease (reprolysin type) with thrombospondin type 1 motif, 5 (aggrecanase-2)	ADAMTS5	W73846	1.4	6.1	2.8
69622_at			AI669987	5.5	6.1	4
63331_at			R74561	6.9	6.1	6.7
38684_at	ATPase, Ca++ transporting, type 2C, member 1	ATP2C1	AJ010953	5.2	6	2.2
39434_at	KIAA0592 protein		AB011164	1.5	6	0.6
1638_at	ets variant gene 6 (TEL oncogene)	ETV6	U11732	4.2	6	1.6

Figure 15EE

41592_at	JAK binding protein			AB000734	4.9	6	7.7
36659_at	collagen, type IV, alpha 2	COL4A2		X05610	1.1	6	2.4
78077_at				A1125204	14.5	6	3
54782_at				A1281068	22.7	6	1.1
50763_at	KIAA1080 protein; Golgi-associated, gamma-adaptin ear containing, ARF-binding protein 2			AA779863	5.1	6	3.4
68165_at	hypothetical protein FLJ10890			AA523442	2.3	6	2.8
86737_at				H88330	10.2	6	4.1
66148_i_at				AW000952	1	6	3.7
64887_at	putative transcription regulation nuclear protein; KIAA1689 protein			AW051603	6.6	6	0.5
54990_at	hypothetical protein			AW009586	16.3	6	8.2
78757_at	selenoprotein X, 1	SEPX1		W92110	3.7	6	0.4
73218_at	phospholipid transfer protein	PLTP		AW007705	8.5	6	1.5
48098_at	glucose regulated protein, 58kD	GRP58		AA478520	2.6	6	6.6
63643_g_at				A1458735	4.6	6	2.2
56648_g_at	KIAA1150 protein			A1968307	2.6	6	0
46079_at				W63785	10	6	5.1
64304_r_at				AW025092	0.6	6	0.8
55061_at				AA522882	10.2	6	1.4
35789_at	KIAA1042 protein			AB028965	6	6	6.4
64269_at				AW023766	4.3	6	8.5
40193_at	enolase 2, (gamma, neuronal)	ENO2		X51956	7.1	5.9	3.5
40773_at	myosin, light polypeptide 5, regulatory	MYL5		L03785	5.6	5.9	0
40615_at	eukaryotic translation initiation factor 3, subunit 1 (alpha, 35kD)	EIF3S1		AA780049	4.1	5.9	0.6
37603_at	interleukin 1 receptor antagonist	IL1RN		X52015	8.9	5.9	5.6
48025_at	NCK adaptor protein 2	NCK2		A1934552	51.7	5.9	6.9
49875_at				A1733353	6.3	5.9	16.6
32156_at	poliovirus receptor-related 2 (herpesvirus entry mediator B)	PVRL2		AF044968	23.4	5.9	8.9
62283_s_at				AA195408	1.6	5.9	0.2
45337_at				W72338	5.6	5.9	1.6
64099_at	hypothetical protein FLJ10700			W81185	6.2	5.9	1.4
44652_at	hypothetical protein FLJ22313			AL048867	4.1	5.9	0
43391_at	porcupine			AW002763	15.7	5.9	1.8
45181_at	zinc finger protein 281	ZNF281		AA121673	5.1	5.9	9.5
54049_at				A1652991	3.8	5.9	0

Figure 15FF

54579_at				AI799534	0.1	5.9	14.6
55055_at				N22687	4	5.9	7.5
62994_at	activating transcription factor 5		ATF5	H50590	7.2	5.9	0.3
86679_at				T79952	1.8	5.9	0.4
56435_at	hypothetical protein FLJ20625			AA573749	6.7	5.9	8.5
56546_at	KIAA1545 protein			H17730	4.9	5.9	5.8
41611_at	thyroid hormone receptor interactor 4		TRIP4	L40371	2.8	5.8	2.9
1294_at	ubiquitin-activating enzyme E1-like		UBE1L	L13852	2.5	5.8	3.2
35938_at	phospholipase A2, group IVA (cytosolic, calcium-dependent)		PLA2G4A	M72393	12.4	5.8	0.2
35844_at	syndecan 4 (amphiglycan, ryudocan)		SDC4	D79206	77.4	5.8	20.7
1183_at	small inducible cytokine subfamily A (Cys-Cys), member 17		SCYA17	D43767	3.2	5.8	1.7
36171_at	activated RNA polymerase II transcription cofactor 4			AI521453	4.2	5.8	2.7
36609_at	solute carrier family 1 (glial high affinity glutamate transporter), member 3		SLC1A3	D26443	34.2	5.8	3.4
36943_r_at	pleiomorphic adenoma gene-like 1		PLAGL1	U81992	8.1	5.8	7.6
38010_at	BCL2/adenovirus E1B 19kD-interacting protein 3		BNIP3	AF002697	19.5	5.8	0.2
38436_at	KIAA0252 protein			D87440	0.2	5.8	7.5
38824_at	Tat-interacting protein (30kD)			AF039103	5.1	5.8	12.6
37068_at	phospholipase A2, group VII (platelet-activating factor acetylhydrolase, plasma)		PLA2G7	U24577	0	5.8	0
45379_at				AI659019	6.4	5.8	9.6
39351_at	CD59 antigen p18-20 (antigen identified by monoclonal antibodies 16.3A5, EJ16, EJ30, EL32 and G344)		CD59	M84349	15.9	5.8	0.4
76600_at				AI631895	0.5	5.8	0.2
59279_at				AI698004	1	5.8	1.3
90038_at	sin3-associated polypeptide, 18kD		SAP18	AW006624	3.5	5.8	5.5
80431_at	ras-like protein			AI674511	3	5.8	0
47969_at				AA631997	0	5.8	3.1
48874_f_at				AI969486	3.7	5.8	4.6
63526_f_at				AI669667	5.4	5.8	7.1
53107_at				N20928	0.1	5.8	0
64412_at	Rho guanine exchange factor (GEF) 11		ARHGEF11	AI090007	6.7	5.8	2.5
45072_r_at				AI033014	3.6	5.8	12.1
75421_at				AA927317	8.6	5.8	1.1
58305_at	KIAA1151 protein			AI003520	38.6	5.8	5.2

Figure 15GG

57722_at									
57785_at	H4 histone family, member H	H4FH							
48767_at	ALEX1 protein								
77040_s_at									
49003_at	NADH dehydrogenase (ubiquinone) 1 beta subcomplex, 10 (22kD, PDSW)	NDUFB10							
49080_at									
39053_at	U4/U6-associated RNA splicing factor								
40408_at	cysteinyl-tRNA synthetase	CARS							
41304_at	neuropilin 2	NRP2							
38899_s_at	hypothetical protein FLJ20693								
41355_at	E2F transcription factor 5, p130-binding	E2F5							
37026_at	core promoter element binding protein	COPEB							
33165_at	testis-specific kinase 2	TESK2							
35230_at	hypothetical protein, clone 24751								
36143_at	caspase 3, apoptosis-related cysteine protease	CASP3							
32541_at	protein phosphatase 3 (formerly 2B), catalytic subunit, gamma isoform (calcineurin A gamma)	PPP3CC							
37640_at	hypoxanthine phosphoribosyltransferase 1 (Lesch-Nyhan syndrome)	HPRT1							
34010_at	hypothetical protein from clones 23549 and 23762								
41010_at	Homer, neuronal immediate early gene, 1B								
87995_at	heterogeneous nuclear ribonucleoprotein U (scaffold attachment factor A)	HNRPU							
91772_at									
48484_at	CGI-111 protein								
46497_at									
82887_at									
73238_at	collagen, type IX, alpha 1	COL9A1							
44718_at									
77572_at	hypothetical protein FLJ20429								
79858_at									
69987_at									
44207_at									
64036_at	hypothetical protein FLJ10716								

Figure 15HH

51105_at	Rho guanine nucleotide exchange factor (GEF) 3	ARHGEF3	N57933	0	5.7	13.3
84347_s_at	c6.1A		AA815282	2.7	5.7	8.4
45655_at	tumor antigen SLP-8p		AL045849	5.4	5.7	2.4
46118_at			AI948463	2.9	5.7	0
55959_at			AA148929	3.7	5.7	4.9
48749_at			D11630	2	5.7	5.3
51672_f_at			AI161358	14.7	5.7	16.9
37472_at	mannosidase, beta A, lysosomal	MANBA	U60337	3.1	5.6	3.5
39857_at	syntaxin 11	STX11	AF044309	12.7	5.6	12.9
40496_at	complement component 1, s subcomponent	C1S	J04080	26.3	5.6	7.9
39424_at	tumor necrosis factor receptor superfamily, member 14 (herpesvirus entry mediator)	TNFRSF14	U70321	0.2	5.6	0
35648_at	KIAA0442 protein		AB007902	0.7	5.6	9.9
35787_at	dynein, cytoplasmic, intermediate polypeptide 2	DNCI2	AI986201	5.6	5.6	0
39165_at	nitrogen fixation cluster-like		U47101	2.2	5.6	1.4
58175_at			AA868339	0.6	5.6	3.3
52591_s_at	glutamate-cysteine ligase, catalytic subunit	GCLC	AI971137	0	5.6	17.5
78537_f_at			F24633	1.6	5.6	0
69874_at	casein kinase 1, gamma 1	CSNK1G1	AI805367	0	5.6	1.2
85691_at			AI989354	5.3	5.6	6.1
45272_at			AA058569	5.2	5.6	2.4
79105_at			AA830599	5	5.6	2.2
88746_at			AI671816	2.3	5.6	2
44183_at			H05653	8.9	5.6	2.2
80084_at			AI889019	16.3	5.6	0.8
65082_at	ficolin (collagen/fibrinogen domain-containing) 3 (Hakata antigen)	FCN3	AI150905	0	5.6	0.6
87860_s_at	solute carrier family 21 (organic anion transporter), member 12	SLC21A12	AW016409	11.9	5.6	0
83558_i_at			AI809148	0.9	5.6	0
68299_at			R07844	7.5	5.6	4.4
41830_at	KIAA0494 gene product		AB007963	9	5.6	6.4
34481_at	vav 1 oncogene	VAV1	AF030227	4.5	5.6	5.9
53708_at	cation-chloride cotransporter-interacting protein		AI467928	6.1	5.6	5.2
49390_at			AA999858	5.6	5.6	5.6
50244_at	nucleotide binding protein 2 (E.coli MinD like)	NUBP2	AA552351	4.8	5.6	6.4

Figure 15II

37148_at	leukocyte immunoglobulin-like receptor, subfamily B (with TM and ITIM domains), member 3	LILRB3	AF025533	7.3	5.5	0.2
39749_at	proteasome (prosome, macropain) 26S subunit, non-ATPase, 4	PSMD4	U51007	6.6	5.5	0
40455_at	KIAA0830 protein		AB020637	1.2	5.5	2.9
32640_at	intercellular adhesion molecule 1 (CD54), human rhinovirus receptor	ICAM1	M24283	19.2	5.5	1.4
36484_at	UDP-N-acetyl-alpha-D-galactosamine:polypeptide N-acetylgalactosaminyltransferase 3 (GalNAc-T3)	GALNT3	AI935146	12.9	5.5	8.4
37948_at	ATPase, H+ transporting, lysosomal (vacuolar proton pump) 42kD	ATP6C	J05682	3.5	5.5	0.1
56350_at			AI928492	6.4	5.5	1.5
66269_at			AA669114	0	5.5	1.5
81276_at			N63894	3.5	5.5	7.2
52848_at	KIAA0792 gene product		AI491983	3.4	5.5	1.8
57617_at			AW009562	1	5.5	1
74975_f_at	zinc finger protein 173	ZNF173	AW009746	5.5	5.5	14.1
50563_r_at			AI224126	3.9	5.5	2.5
43281_at			AA481181	3.5	5.5	7.1
87743_at			AA884171	1.4	5.5	1.7
79360_at			AA555312	3.2	5.5	3.9
76515_at	hypothetical protein DKFZp566G1424		AA777011	11.6	5.5	0
59627_at	hypothetical protein FLJ10628		AA885169	2	5.5	7
58350_at	hypothetical protein FLJ20651		AW014619	3.8	5.5	0.6
45302_at			AL042856	8.7	5.5	0.6
83227_at			AI939517	0.6	5.5	1.6
74456_r_at			AI970474	6.5	5.5	4
45661_at			N90348	4.5	5.5	6.7
73146_at	vascular endothelial growth factor	VEGF	AW024572	12.9	5.5	1.6
46601_at	ubiquinol-cytochrome c reductase hinge protein	UQCRRH	T16388	1.8	5.5	0
62248_at	metallothionein 1E (functional)	MT1E	R80823	10	5.5	3.1
31680_at			M55630	14.3	5.5	6.7
1252_at	DNA segment, single copy probe LNS-CAI/LNS-CAII (deleted in polyposis)		M73547	6.9	5.5	6.6
35310_at			D45288	9.4	5.5	8.9
45521_s_at	KIAA0100 gene product		AI967955	5.4	5.5	5.7
55525_at			AI972498	6.2	5.5	6.3

Figure 15JJ

38971_r_at	Nef-associated factor 1			AJ011896	18.5	5.4	1.5
38571_at	FGFR1 oncogene partner			Y18046	4.2	5.4	0
32227_at	proteoglycan 1, secretory granule	PRG1		X17042	7.5	5.4	4.5
40555_at	phosphatidylinositol glycan, class F	PIGF		AL043108	4.2	5.4	0.2
1046_at	retinoic acid- and interferon-inducible protein (58kD)			U34605	7	5.4	3
36971_at	KIAA0257 protein			D87446	4.8	5.4	2
41532_at	zinc finger protein 151 (pHZ-67)	ZNF151		Y09723	13.8	5.4	4.2
37025_at	LPS-induced TNF-alpha factor			AL120815	14.1	5.4	0.7
37351_at	uridine phosphorylase	UP		X90858	6.2	5.4	0.2
37890_at	CD47 antigen (Rh-related antigen, integrin-associated signal transducer)	CD47		X69398	3.7	5.4	1
33134_at	adenylate cyclase 3	ADCY3		AB011083	1.6	5.4	0
74988_at				AI074017	3.4	5.4	3
42428_r_at				AA463189	6.7	5.4	6.4
59887_r_at				AI733573	2.4	5.4	8.8
73152_s_at				AI138359	11.5	5.4	0
48740_s_at	novel retinal pigment epithelial gene			AI973108	18.7	5.4	4.1
55037_r_at				AI220443	4.1	5.4	6.4
43850_s_at	PP1201 protein			AA668732	3.4	5.4	5
45306_at				AA742560	1.2	5.4	13.2
58411_at	KIAA1610 protein			AL037424	7.4	5.4	12.2
71661_at				AI174651	9.1	5.4	6.5
87221_at				AA829405	3.5	5.4	3.1
43576_at	hypothetical protein			AI348427	0.6	5.4	5.9
46105_at				AA149641	5.4	5.4	0
1526_i_at	fibroblast growth factor 8 (androgen-induced)	FGF8		U47011	0	5.4	0.4
47318_at				AI949175	29.7	5.4	1.3
44376_at	KIAA0582 protein			AA973070	12.9	5.4	7.3
88064_at				AA543076	2.6	5.4	0.8
49053_at				AA470002	4.5	5.4	4.8
85959_at				AI016237	3.2	5.4	2.6
70532_r_at				AA019714	6.8	5.4	3.2
40137_at	protein tyrosine phosphatase, non-receptor type 1	PTPN1		M31724	12.9	5.3	5.2
411_i_at	interferon induced transmembrane protein 2 (1-8D)	IFITM2		X57351	0.8	5.3	2
41156_g_at	catenin (cadherin-associated protein), alpha 1 (102kD)	CTNNA1		U03100	9.1	5.3	3.7

Figure 15KK

34258_at	muskelin 1, intracellular mediator containing kelch motifs	MKLN1	W28205	6.9	5.3	0.7
39929_at	KIAA0922 protein		AB023139	1.8	5.3	1.7
34409_at	DKFZP564C1940 protein		AL080164	7.7	5.3	0
36991_at	splicing factor, arginine/serine-rich 4	SFRS4	L14076	5.4	5.3	4.2
43459_at	KIAA0680 gene product		AI472209	0	5.3	5.1
65130_at			AI749588	0.1	5.3	0
81317_at			AI345709	8.2	5.3	0.8
85035_i.at			H75345	2.8	5.3	0.9
67989_at			AA825836	6	5.3	1.2
85724_at			AI634355	15.8	5.3	2.2
63439_at			AA249119	1.9	5.3	3
59660_at			AI800240	12.2	5.3	1.2
70319_at			AI766604	3.3	5.3	0
44089_at			W26293	2.1	5.3	3.7
42363_r.at	STAT induced STAT inhibitor 3		AI680350	41.7	5.3	5.8
74324_r.at			AA724785	2.8	5.3	1.9
53164_at			AI796566	9	5.3	0.1
53100_at	hypothetical protein FLJ10307		AA747448	34.8	5.3	1.3
58622_at			AA463374	3.5	5.3	6
51883_at	hypothetical protein		AI972120	4.1	5.3	0
69007_r.at			AI583317	1.8	5.3	0.8
69970_at			AI086392	5.1	5.3	7.2
72520_at			AA161305	1.6	5.3	0
55567_at			AA147933	5.3	5.3	5.2
58954_at	NADH dehydrogenase (ubiquinone) Fe-S protein 8 (23kD) (NADH-coenzyme Q reductase)	NDUFS8	AW021179	6.4	5.3	6.5
1787_at	cyclin-dependent kinase inhibitor 1C (p57, Kip2)	CDKN1C	U22398	0.3	5.2	45.6
41430_at	WD repeat domain 7	WDR7	AB011113	0	5.2	0
37466_at	RAB7, member RAS oncogene family-like 1	RAB7L1	D84488	11.9	5.2	1.2
31867_at			AF052174	1.8	5.2	3.6
31883_at	5-methyltetrahydrofolate-homocysteine methyltransferase reductase	MTRR	AF025794	16.2	5.2	0
34131_at	solute carrier family 7, (cationic amino acid transporter, y+ system) member 11	SLC7A11	AB026891	7	5.2	2.4
38977_at	tyrosyl-tRNA synthetase	YARS	U89436	11.4	5.2	0.6

Figure 15LL

32776_at	v-ral simian leukemia viral oncogene homolog B (ras related; GTP binding protein)	RALB	M35416	11.3	5.2	6.4
40639_at	SCO (cytochrome oxidase deficient, yeast) homolog 2	SCO2	AL021683	7.1	5.2	30.4
38332_at	purinergic receptor P2X, ligand-gated ion channel, 4	P2RX4	U83993	6.3	5.2	0.5
34311_at	glutaredoxin (thioltransferase)	GLRX	X76648	1.6	5.2	1.1
39982_r_at	macrophage scavenger receptor 1	MSR1	D13265	0	5.2	0.6
37188_at	phosphoenolpyruvate carboxykinase 2 (mitochondrial)	PCK2	X92720	4.3	5.2	0
36977_at	N-ethylmaleimide-sensitive factor attachment protein, alpha	NAPA	U39412	2.7	5.2	2.9
40390_at	serine dehydratase	SDS	J05037	3.7	5.2	16.9
38114_at	RAD21 (S. pombe) homolog	RAD21	D38551	7.4	5.2	5.1
276_at	DnaJ (Hsp40) homolog, subfamily A, member 1	DNAJA1	L08069	8.2	5.2	6
48014_at			N54957	38.8	5.2	2
54747_at	endosulfine alpha	ENSA	AA167734	5.1	5.2	0
91799_at			AA586814	5.4	5.2	0
47951_at			AA521145	5.4	5.2	5.2
74864_at	hypothetical protein		H94320	7.7	5.2	2.5
51607_at	KIAA1145 protein		H06311	9.3	5.2	2.8
65814_at			N95618	5.5	5.2	4.5
65531_at	hypothetical protein FLJ22794		AA948676	1.9	5.2	10.5
32855_at	low density lipoprotein receptor (familial hypercholesterolemia)	LDLR	L00352	9.6	5.2	0.2
72723_at			A1754075	6.1	5.2	3.1
44792_s_at			AA972102	0.8	5.2	3.2
49508_at			A1041540	9.1	5.2	3.5
82032_at			A1823999	4.6	5.2	3.5
53920_at			A1655289	30.5	5.2	0.7
55940_at			A1701480	0	5.2	21.4
89908_f_at	GDP dissociation inhibitor 1	GDI1	AW004027	3.5	5.2	1
51161_at	ras homolog gene family, member A	ARHA	A1743085	20.9	5.2	20.3
1203_at			D17357	17.8	5.2	10.3
32842_at	B-cell CLL/lymphoma 7A	BCL7A	X89984	7.4	5.2	6.1
44080_at			W68180	6.3	5.2	5.5
45736_at	two-pore channel 1, homolog		AL048491	5.4	5.2	5.1
64330_s_at			AW002183	6	5.2	6.6
54152_at	eukaryotic translation initiation factor 4E binding protein 1	EIF4EBP1	A1026669	5.8	5.2	5.8
35064_at			X81006	6.4	5.1	10.5

Figure 15MM

38968_at	SH3-domain binding protein 5 (BTK-associated)	SH3BP5	AB005047	2.1	5.1	1
39852_at	KIAA0610 protein		AB011182	2.2	5.1	7.4
38129_at	glycerol kinase	GK	L13943	3.5	5.1	1.7
40063_at	nuclear domain 10 protein		U22897	7.9	5.1	9.3
40781_at	serologically defined colon cancer antigen 8	SDCCAG8	U79271	0.7	5.1	1.5
40922_at			AF010235	4.6	5.1	2.2
1061_at	interleukin 10 receptor, alpha	IL10RA	U00672	9.6	5.1	11.1
41585_at	KIAA0746 protein		AB018289	1.9	5.1	3.1
36535_at	microfibrillar-associated protein 1	MFAP1	U04209	5.8	5.1	7.1
36845_at	KIAA0136 protein		D50926	7.2	5.1	5.4
40364_at	solute carrier family 31 (copper transporters), member 1	SLC31A1	U83460	0	5.1	0.8
36678_at	transgelin 2	TAGLN2	D21261	9.2	5.1	2.4
57737_at			A1150441	3.2	5.1	3.4
58455_at			A1735252	0	5.1	2.8
56894_at	RAB38, member RAS oncogene family	RAB38	N29070	2.6	5.1	6.5
67329_at	hypothetical protein FLJ22833		AA610377	13.3	5.1	0.8
57675_at			AW052119	7.8	5.1	3.3
80603_at			A1264334	4	5.1	1
75046_at	HS1 binding protein		A1970711	6.7	5.1	1.3
47239_r_at			A1768586	1.5	5.1	4.4
44742_s_at	hypothetical protein FLJ21939 similar to 5-azacytidine induced gene 2		AA205994	10.7	5.1	4.7
45270_at	CGI-127 protein		AA151651	2.8	5.1	0.8
68748_at			A1651275	0	5.1	1.8
44375_at			R40515	0.6	5.1	1.4
53659_r_at	DKFZP434D193 protein		AA453704	3.5	5.1	1
50205_at	leptin receptor	LEPR	N46838	5.2	5.1	15
43056_at			AA780061	10.4	5.1	0
65776_at			A1341468	11.3	5.1	0
42804_at			W86160	3.7	5.1	5.4
43438_at			A1694413	0.5	5.1	7.4
54830_r_at			A1203373	1.1	5.1	1.3
45110_at			A1989567	0	5.1	0.2
46337_at			AA188763	2.2	5.1	3
64769_at			A1298599	16.8	5.1	2.9

Figure 15NN

58341_i_at			AI734899	3.7	5.1	1.6
43445_at			AI123567	2.8	5.1	0.5
47165_at	small nuclear ribonucleoprotein polypeptide G	SNRPG	AI582592	8.5	5.1	2
50204_at			AA456955	2.6	5.1	2.5
46722_at	cathepsin D (lysosomal aspartyl protease)	CTSD	AI807430	9.6	5.1	0
49709_at	hypothetical protein FLJ10116		AA127924	3.5	5.1	0.4
43361_s_at	syndecan 1	SDC1	AI972735	3.7	5.1	0
53806_at	hypothetical protein		W22715	2	5.1	2.4
40585_at	adenylate cyclase 7	ADCY7	D25538	7.2	5.1	7.2
36222_at	dimethylarginine dimethylaminohydrolase 2	DDAH2	AJ012008	4.1	5.1	0.1
60061_at	arachidonate 5-lipoxygenase-activating protein	ALOX5AP	AI983204	5.5	5.1	5.4
44078_at			W95017	4.1	5.1	5.5
61853_at			AA773816	5.1	5.1	5.1
53881_s_at			AL044098	6.1	5.1	5.8
59136_at			AA779895	5.2	5.1	5.2
46664_at	apoptosis-associated speck-like protein containing a CARD		AI148558	5.7	5.1	6.3
37442_at			AL050378	27.6	5	11.8
41620_at	KIAA07116 gene product		AB018259	8.9	5	0
39412_at	zinc finger protein 173	ZNF173	U09825	4.3	5	16.9
32645_at	KIAA0477 gene product		AB007946	4.2	5	4.6
39432_at	UDP-Gal:betaGlcNAc beta 1,4- galactosyltransferase, polypeptide 4	B4GALT4	AF038662	8.1	5	1.3
31832_at	KIAA0286 protein		AB006624	41.3	5	5.1
222_at	exostoses (multiple) 1	EXT1	S79639	50.9	5	9.8
33697_at	purinergic receptor P2X, ligand-gated ion channel, 7	P2RX7	Y12851	9.1	5	0.9
33847_s_at	cyclin-dependent kinase inhibitor 1B (p27, Kip1)	CDKN1B	AI304854	9.9	5	8.2
32202_at	HBV associated factor		U67322	3.3	5	2.8
35145_at	MAX binding protein	MNT	X96401	4.8	5	0
35238_at	TNF receptor-associated factor 5	TRAF5	AB000509	17	5	0
36827_at	golgi resident protein GCP60		AF020762	7.3	5	2.7
36780_at	clusterin (complement lysis inhibitor, SP.40,40, sulfated glycoprotein 2, testosterone-repressed prostate message 2, apolipoprotein J)	CLU	M25915	32.9	5	8.7
237_s_at	protein phosphatase 2 (formerly 2A), catalytic subunit, alpha isoform	PPP2CA	M60483	7.6	5	8.6
34041_at	small inducible cytokine subfamily A (Cys-Cys), member 22	SCYA22	U83171	6.9	5	5.4
63316_at	RNA binding motif protein 8B	RBM8B	AL047586	6.5	5	0

Figure 1500

75747_at			AA626334	2.3	5	4
66288_at			AI769560	2	5	2.6
66326_at			AI936888	9.5	5	3.8
81410_at			AI830490	9	5	1
53718_at	hepatocellular carcinoma-associated antigen 112		AI684645	0.6	5	0.1
73850_at			AI084934	6.5	5	6.5
53799_at	hypothetical protein similar to mouse DnaI1		AA628434	28.3	5	0
49515_at			AI041860	1.7	5	2.1
78054_at			AI206947	2.2	5	0.6
54064_i_at			AA903287	14	5	5.7
48357_at	ribosomal protein S28	RPS28	AA968694	0.2	5	18.1
75764_at			AA808948	4.2	5	0.4
704_at	nuclear factor I/A	NFIA	U07809	3.8	5	5
89316_at			AI809752	1.3	5	5.1
51900_g_at	hypothetical protein FLJ10120		AA987531	0.6	5	3.5
52204_at			AA894564	4.7	5	4.3
53128_at	hypothetical protein FLJ10494		AI678076	6.3	5	3.2
44099_at			N34505	2.9	5	1.4
62801_at	KIAA0554 protein		AA937109	6.8	5	1.1
45893_at			H24387	7.5	5	7.6
52042_at			AA081395	7.9	5	0
53661_at			AL038450	16.2	5	5.2
45055_at			AA461205	7.8	5	5.5
47403_at			AI990290	2	5	1
46215_at			AL079949	7.9	5	2.5
41950_at			AA058314	0.7	5	0
46733_at			AI340046	1.2	5	3
54098_at	hypothetical protein FLJ10201		AI816255	7.1	5	1.5
65715_at	nuclear receptor coactivator 1	NCOA1	AL079440	2.5	5	6.7
55063_at	glutamate decarboxylase 1 (brain, 67kD)	GAD1	AL042399	31.4	5	0
70212_at			AI769797	7.8	5	0.4
68028_at			AI383877	1.5	5	1
50417_at			AA641023	6.8	5	6.6
51123_at			H11724	5.2	5	4.9
64879_at	heparan sulfate 6-O-sulfotransferase	HS6ST	AA206625	3	5	4.3

Figure 15PP

40960_at	UDP-Gal:betaGlcNAc beta 1,4-galactosyltransferase, polypeptide 1	B4GALT1	D29805	16.7	4.9	3.9
41488_at	hypothetical protein A-211C6.1		AC002394	8.2	4.9	4
34308_at	H2A histone family, member L	H2AFL	U90551	14	4.9	13.2
36578_at	baculoviral IAP repeat-containing 2	BIRC2	U37547	18.6	4.9	13.5
40357_at	inhibin, beta A (activin A, activin AB alpha polypeptide)	INHBA	J03634	38.6	4.9	5
36193_at	partner of RAC1 (arfaptin 2)		U52522	2.8	4.9	0
1461_at	nuclear factor of kappa light polypeptide gene enhancer in B-cells inhibitor, alpha	NFKBIA	M69043	17	4.9	6.3
37926_at	Kruppel-like factor 5 (intestinal)	KLF5	D14520	2.1	4.9	2.2
58192_at	KIAA1411 protein		AI308839	6	4.9	3.7
54961_r_at			AA203620	10.9	4.9	1.1
47818_at			AI968249	2.4	4.9	6.9
46529_at			AA988769	1.5	4.9	3.5
90148_at	putative ATPase		AI694453	0	4.9	0.1
80361_at			AI268476	7.8	4.9	1
83245_at			AI300111	7.7	4.9	9.2
67087_at			AI351909	10.3	4.9	4.9
65188_at			AA418143	3.3	4.9	1.7
84145_at	hypothetical protein FLJ23360		AA907658	3.3	4.9	3.1
51183_at	transcriptional intermediary factor 1 gamma		N98658	7.5	4.9	4.9
58916_at			AI672101	0	4.9	2.7
83419_at			AI078033	2.8	4.9	2.9
64623_at	CDC10 (cell division cycle 10, S. cerevisiae, homolog)	CDC10	AA015969	3.7	4.9	2.4
87391_at			AA707322	5.4	4.9	0.3
48390_at			AA430602	4.6	4.9	6.3
63908_r_at			R46086	2	4.9	3.4
80429_at			AI796486	2.1	4.9	0.2
89128_at	C-type (calcium dependent, carbohydrate-recognition domain) lectin, superfamily member 9	CLEC5F9	AI968491	7.5	4.9	6.6
73002_at	matrix metalloproteinase 14 (membrane-inserted)	MMP14	AI052526	11.4	4.9	0
50413_at			AA418534	1.1	4.9	2.1
75899_r_at			AA677461	0.3	4.9	2.9
88550_at	hypothetical protein FLJ12929		AI468004	23.6	4.9	0
44031_at			AI478812	0.9	4.9	0
63662_at			AI126468	12.6	4.9	4.5

Figure 15QQ

45143_at				AA160973	15.8	4.9	0.2
51984_at				AA148813	9.7	4.9	0.3
41992_at				AI002966	3.1	4.9	24.9
54989_at				H43374	4.8	4.9	1.2
48315_at				AA286940	4.2	4.9	5.5
72603_at				AI620888	3.6	4.9	3.1
32550_r_at	CCAAT/enhancer binding protein (C/EBP), alpha		CEBPA	Y11525	5.9	4.9	4.6
53857_at	hypothetical protein HES6			H09392	1.8	4.9	5.1
49633_at	MADS box transcription enhancer factor 2, polypeptide D (myocyte enhancer factor 2D)		MEF2D	N21131	4.9	4.9	4.8
35434_at				L16794	1.1	4.8	4.6
37451_at				AL109695	50.6	4.8	5.8
39777_at	KIAA0916 protein			AF075587	0.4	4.8	6.3
32643_at	glucan (1,4-alpha-), branching enzyme 1 (glycogen branching enzyme, Andersen disease, glycogen storage disease type IV)		GBE1	L07956	4.1	4.8	0.4
40898_at	sequestosome 1		SQSTM1	U46751	2	4.8	3.3
32150_at	golgi autoantigen, golgin subfamily a, 4		GOLGA4	X82834	13.2	4.8	2.2
1323_at	ubiquitin B		UBB	X04803	9.2	4.8	3.4
287_at	activating transcription factor 3		ATF3	L19871	9.2	4.8	15.2
36060_at	signal recognition particle 54kD		SRP54	U51920	1.5	4.8	1.8
36899_at	special AT-rich sequence binding protein 1 (binds to nuclear matrix/scaffold-associating DNA's)		SATB1	M97287	4.3	4.8	6.6
37001_at	calpain 2, (m/II) large subunit		CAPN2	M23254	4.1	4.8	2.6
38778_at	KIAA1046 protein			AB028969	5.3	4.8	3.3
59617_at	RNA polymerase I transcription factor RRN3			AA830682	0	4.8	0
76118_at	hypothetical protein FLJ22655			AI498375	4.2	4.8	0.8
90975_i_at				N22605	6.6	4.8	0.9
60906_r_at	hypothetical protein FLJ10773			AA044181	4.4	4.8	3.9
83491_at				AI347066	1.2	4.8	2.6
59000_at	x 010 protein		x 010	AA629304	6.6	4.8	4
84068_at				AI375097	3.1	4.8	3.7
62334_s_at				T94015	3.9	4.8	0.8
59766_at				AA904413	1.3	4.8	0
54826_at				AA284268	7.8	4.8	3.3
68194_at				AA291194	3.1	4.8	10

Figure 15RR

63868_at	p300/CBP-associated factor	PCAF	AW021507	0.3	4.8	2
49498_at	small protein effector 1 of Cdc42		H10816	9.7	4.8	4.9
44920_at			A1914925	12.7	4.8	7.6
33374_at	complement component 2	C2	L09708	6	4.8	1.7
57270_f_at	glutaminase	GLS	A1968420	4.5	4.8	0.3
57670_at	phosphoprotein associated with GEMs		A1860212	13.5	4.8	1.2
42933_at			T95101	15.3	4.8	11.4
63373_at			AA194033	4.8	4.8	2.2
87415_at			W73083	22.7	4.8	17.3
56242_at	KIAA0493 protein		AA863228	3.8	4.8	2.5
79842_at			A1688250	6.9	4.8	2.4
65837_at	hypothetical protein MGC5306		W93232	2.3	4.8	6.4
43533_s_at	MAGEF1 protein		A1889519	9	4.8	2.6
63368_at			W81119	0	4.8	22
51053_at			A122787	3.8	4.8	6.7
57085_s_at	DKFZP586G0522 protein		AW008790	85.4	4.8	52.3
36929_at	laminin, beta 3 (nicein (125kD), kalinin (140kD), BM600 (125kD))	LAMB3	U17760	10.9	4.8	1.7
47091_at			W86630	1.1	4.8	2
55558_at			AA524361	4.9	4.8	3.6
44868_s_at			AA917734	11.1	4.8	1.9
47544_at			AA741324	22.1	4.8	5.3
46700_at			W55956	2.6	4.8	0.6
42204_at			A1219461	70.7	4.8	5
40113_at	GS3955 protein		D87119	0	4.7	13
32083_at	transmembrane 7 superfamily member 1 (upregulated in kidney)	TM7SF1	AF027826	7.1	4.7	0.4
36372_at	hexokinase 3 (white cell)	HK3	U51333	0	4.7	0
40517_at	KIAA0372 gene product		AB002370	2	4.7	0
33812_at			AL049415	10.7	4.7	0.1
37493_at			H04668	4.6	4.7	3.3
34837_at	KIAA0376 protein		AB002374	1.6	4.7	0
36502_at	PFTAIRE protein kinase 1	PFTK1	AB020641	0.9	4.7	1.1
35365_at	integrin-linked kinase	ILK	U40282	1	4.7	3.7
1237_at	immediate early response 3	IER3	S81914	20.8	4.7	0
37245_at	GA-binding protein transcription factor, alpha subunit (60kD)	GABPA	U13044	2.7	4.7	1.4
56401_at	zinc finger protein 22 (KOX 15)	ZNF22	A1951311	7.8	4.7	6.3

Figure 15SS

44420_at	von Willebrand factor	VWF	R91271	2.7	4.7	1.1
75801_at			A1620827	2	4.7	0
73703_at			AA777616	7	4.7	7.5
76710_at	cardiotrophin-like cytokine; neurotrophin-1/B-cell stimulating factor-3		A1040033	92.7	4.7	4
79074_at			A1524250	2.3	4.7	0
58309_at			A1802060	2.6	4.7	3.4
67505_at			A1435586	5.8	4.7	1
65491_at	hypothetical protein FLJ11085		AL046038	2	4.7	0.9
83065_at			A1638449	3.9	4.7	3.1
49306_at	FE65-LIKE 2		A1890191	6.7	4.7	0.2
45258_at	hypothetical protein DKFZp761G2113		AF039240	9.2	4.7	3.2
54822_at	ubiquitin specific protease 25	USP25	A1672446	1.5	4.7	1.8
67963_at			AA436007	4.7	4.7	3.1
87646_at			AA677692	3.2	4.7	1.6
69448_i_at			A1700454	2.8	4.7	4.6
49626_at	Alg5, S. cerevisiae, homolog of		AA425251	18.6	4.7	3.8
48218_at			T86283	0.5	4.7	1.9
51220_at	small EDRK-rich factor 1A (telomeric)	SERF1A	W67895	13.1	4.7	3.6
81806_at	KIAA1681 protein		A1694702	3.7	4.7	4.6
45608_at			A1202327	11.7	4.7	7.2
60887_s_at			A1950468	2.9	4.7	2.1
47379_at			AA024696	2.5	4.7	2.1
42513_at	ATPase, Na+/K+ transporting, beta 3 polypeptide	ATP1B3	AA151771	8.1	4.7	6.4
55095_at	hypothetical protein		A1279547	1.2	4.7	0
73464_r_at			AA648038	2.2	4.7	2.5
86322_r_at			T53160	1.7	4.7	0.3
40382_at	corticotropin releasing hormone	CRH	V00571	5	4.7	3.8
32562_at	endoglin (Osler-Rendu-Weber syndrome 1)	ENG	X72012	4.6	4.7	6
52261_at			A1970054	5.6	4.7	6.4
38630_at			AL080192	4.7	4.6	1.8
41438_at	KIAA1451 protein		AL049923	3.3	4.6	0.4
1313_at	proteasome (prosome, macropain) subunit, beta type, 7	PSMB7	D38048	3.4	4.6	0.5
41634_at	KIAA0256 gene product		D87445	5.3	4.6	2.4
39703_at	ubiquitously transcribed tetratricopeptide repeat gene, X chromosome	UTX	AF000993	10.2	4.6	48.9

Figure 15TT

960_g_at	cell division cycle 42 (GTP-binding protein, 25kD)	CDC42	M35543	9.8	4.6	2.4
2066_at	BCL2-associated X protein	BAX	L22474	4.1	4.6	9.8
41183_at	cleavage stimulation factor, 3' pre-RNA, subunit 3, 77kD	CSTF3	U15782	5.9	4.6	6.3
33342_at	RNA, U transporter 1	RNUT1	AF039029	0.4	4.6	0.5
31671_at	RNA binding motif, single stranded interacting protein 1, pseudogene	RBMS1P	D82351	3.7	4.6	3.9
473_g_at	signal transducer and activator of transcription 5B	STAT5B	U48730	12.6	4.6	1.9
35198_at	hypothetical protein from clone 24796		AF070596	5.6	4.6	2.4
32304_at			AF035594	0.8	4.6	2
40370_f_at	HLA-G histocompatibility antigen, class I, G	HLA-G	M90683	7	4.6	1.9
36130_f_at	metallothionein 1E (functional)	MT1E	R92331	78.7	4.6	8.6
37221_at	protein kinase, cAMP-dependent, regulatory, type II, beta	PRKAR2B	M31158	6	4.6	5
37732_at	RING1 and YY1 binding protein	RYBP	AL049940	15.1	4.6	5.3
37914_at	KIAA0305 gene product		AB002303	0	4.6	0
38817_at	sperm associated antigen 7	SPAG7	AF047437	3.4	4.6	1
74134_at			AI190292	2.9	4.6	0
89166_at			AI953998	4.8	4.6	0.7
48526_at	hypothetical protein FLJ10582		AI190311	3.6	4.6	1.1
71008_at			AI863088	1.9	4.6	4.1
71035_at			AW003971	1.1	4.6	0.4
57131_at			AA419427	4.1	4.6	10.4
51121_at			R45590	8.8	4.6	5.2
50081_at			AI765698	3.2	4.6	3.4
50323_g_at	hypothetical protein FLJ10143		AI656132	3.3	4.6	0.2
91206_at			AI683911	3.7	4.6	0.3
56067_at	centrosomal P4.1-associated protein; uncharacterized bone marrow protein BM032		AI288949	1.6	4.6	2.5
69284_at			AI655913	1.5	4.6	0.2
45922_at			AI560775	2.6	4.6	8.4
62561_at	DnaJ (Hsp40) homolog, subfamily B, member 4	DNAJB4	AI741503	18.5	4.6	26.4
57579_at			AI680721	14	4.6	4.6
54479_at			AA214559	0	4.6	0
53672_at			AA055179	2.2	4.6	11.8
62557_at	hypothetical protein FLJ12820		AI745595	3.8	4.6	0.8
62993_at			AA521445	0.5	4.6	0

Figure 15UU

61440_at			AA448535	0	4.6	0
42093_at			AA398660	3.5	4.6	4.5
82136_i_at			AA876935	1.9	4.6	2.7
36601_at	vinculin	VCL	M33308	5.2	4.6	5.1
48750_at	class I cytokine receptor		AI983115	5.2	4.6	4.9
43929_at			AA225263	4.7	4.6	4.4
41386_i_at	KIAA0346 protein		AB002344	16.8	4.5	0.5
34982_at	cytochrome P450, subfamily XXVIA, polypeptide 1	CYP26A1	AF005418	2.2	4.5	3.6
975_at	serine/threonine kinase 18	STK18	Y13115	9.5	4.5	12.9
32963_s_at	hypothetical GTP-binding protein DKFZp761H171		W27549	0	4.5	0
1891_at	mitogen-activated protein kinase kinase 8	MAP3K8	D14497	53.8	4.5	11.2
35136_at	hypothetical protein P15-2		AL031387	8.2	4.5	9.7
35228_at	carnitine palmitoyltransferase I, muscle	CPT1B	Y08682	2.5	4.5	4.3
35625_at	CD97 antigen	CD97	X94630	6.1	4.5	0.8
34824_at	ubiquitin 2	UBQLN2	AB015344	10.3	4.5	0.1
34857_at			Z24724	5.7	4.5	0.2
36118_at	nuclear receptor coactivator 1	NCOA1	AJ000882	3	4.5	4.4
32434_at	myristoylated alanine-rich protein kinase C substrate (MARCKS, 80K-L)	MACS	D10522	8.8	4.5	1.6
32574_at	sphingomyelin phosphodiesterase 1, acid lysosomal (acid sphingomyelinase)	SMPD1	X59960	0	4.5	0.1
40191_s_at	KIAA0582 protein		AI761647	6.3	4.5	0
56927_at			W49628	4.3	4.5	7.8
59314_at	retinoic acid induced 1	RAI1	AI926671	5.7	4.5	3.2
42443_at			AA608577	8.5	4.5	5.3
52290_g_at	long-chain fatty acid coenzyme A ligase 5		AA514342	11.7	4.5	0
73648_s_at			AA758527	3.5	4.5	0.6
76721_at			AI253579	1.5	4.5	0.7
89887_s_at	acyl-Coenzyme A oxidase 3, pristanoyl	ACOX3	AI992213	4.2	4.5	0
83917_at			AI377418	3.1	4.5	2.1
90622_at			AI632015	0.7	4.5	2.3
61840_s_at	catenin (cadherin-associated protein), delta 1	CTNND1	AI863000	5.3	4.5	9.9
82669_i_at			AA059369	0.8	4.5	1.4
65480_at			AA166751	4.6	4.5	1.7
62744_at	guanine nucleotide binding protein beta subunit 4		AI857603	1	4.5	4.2

Figure 15VV

51728_at	protease, cysteine, 1 (legumain)	PRSC1	AA910861	1	4.5	2.6
67470_at			AI928355	7.8	4.5	6.4
69446_at			AA862900	4.6	4.5	2.7
42866_s_at			AW003230	1.5	4.5	4.7
51144_at	hypothetical protein FLJ20337		AI952855	3.6	4.5	2.9
71727_r_at			AA262436	3.3	4.5	3.9
64369_s_at			AA219354	0.7	4.5	0
49967_at			AI658816	0.4	4.5	0
91341_at			AI291659	5.9	4.5	3.8
65639_at			AA195801	1.6	4.5	2.4
43317_at			AA127598	4	4.5	1.9
89100_at	KIAA0771 protein		AI733804	6.8	4.5	1.1
46678_at			AI734922	1.7	4.5	6.3
54147_at	pyruvate dehydrogenase kinase, isoenzyme 4	PDK4	AI763378	38.3	4.5	0.6
58332_at			AA399394	1.8	4.5	4.1
43213_r_at			H21152	0	4.5	5.9
83455_r_at			AA644692	2	4.5	0.1
40045_g_at	chromosome 18 open reading frame 1	C18ORF1	AF009425	5.6	4.5	3.1
65171_at	clone FLB4739		N36909	5.1	4.5	5.2
58468_at			C17704	4.5	4.5	4.3
55695_at	glycine cleavage system protein H (aminomethyl carrier)	GCSH	AI890917	5.1	4.5	4.2
41697_at	acyl-Coenzyme A oxidase 3, pristanoyl	ACOX3	Y11411	4.5	4.4	1
32129_at	hypothetical protein, similar to (U06944) PRAJA1		AL079314	6.9	4.4	1.4
40606_at	ELL-RELATED RNA POLYMERASE II, ELONGATION FACTOR		U88629	21.6	4.4	2.6
34199_at	SEC24 (S. cerevisiae) related gene family, member A	SEC24A	AJ131244	2.5	4.4	1.3
32954_at	DKFZP434D193 protein		U79263	3.9	4.4	1
33886_at	spectrin SH3 domain binding protein 1	SSH3BP1	AF006516	12.4	4.4	6.4
38617_at	LIM domain kinase 2	LIMK2	D45906	27.3	4.4	4
41795_at	NCK adaptor protein 1	NCK1	X17576	11.3	4.4	2
35320_at	solute carrier family 11 (proton-coupled divalent metal ion transporters), member 2	SLC11A2	AB004857	5.6	4.4	0
36131_at	dimethylarginine dimethylaminohydrolase 2	DDAH2	AJ012008	1.5	4.4	3.3
36651_at	acid phosphatase 2, lysosomal	ACP2	X15525	2.4	4.4	1.7
37295_at	general transcription factor IIE, polypeptide 2 (beta subunit, 34kD)	GTF2E2	X63469	1.7	4.4	0.2

Figure 15WW

38037_at	diphtheria toxin receptor (heparin-binding epidermal growth factor-like growth factor)	DTR	M60278	3.3	4.4	1.3
38805_at	TG-interacting factor (TALE family homeobox)	TGIF	X89750	6.2	4.4	6.6
38823_s_at	serine/threonine kinase 17a (apoptosis-inducing)	STK17A	AI961743	13.7	4.4	0.3
48055_at			N66640	0	4.4	0
51174_at	spinocerebellar ataxia 7 (olivopontocerebellar atrophy with retinal degeneration)	SCA7	AL043992	3.9	4.4	3.2
48837_at	40S ribosomal protein S27 isoform		AI149884	1.1	4.4	1.9
49684_g_at	splicing factor, arginine/serine-rich 11	SFRS11	AI306386	0.7	4.4	0.9
61896_at			AL038511	3.6	4.4	8.7
76997_at			AI733141	3.2	4.4	1.6
78605_at			AI080388	9	4.4	5.6
61240_at			W27252	2.8	4.4	8.9
82306_r_at			AI827008	3.7	4.4	3.1
45233_at	mesenchymal stem cell protein DSC92		W22743	8.9	4.4	3.6
67764_r_at			AA004879	29	4.4	11.8
87874_at			T69727	4.7	4.4	8.2
88154_r_at			AI248758	1.2	4.4	3.8
48463_at			AA362254	2.5	4.4	3.8
60867_at			N38839	2.5	4.4	1.6
43961_s_at	hypothetical GTP-binding protein DKFZp761H171		AI627648	0.3	4.4	0
55136_r_at			AI822128	2.9	4.4	3.3
65839_at	PC3-96 protein		AW022836	3.7	4.4	6.3
91274_at	chromatin accessibility complex 1	CHRA1	AA988299	4.5	4.4	0.6
53263_r_at			H15157	5.8	4.4	2.6
56308_at	hypothetical protein FLJ23399		AA193501	7.2	4.4	0.2
44556_at	KIAA1344 protein		AA113931	0	4.4	0.4
64457_at	leucine rich repeat (in FLII) interacting protein 2	LRRFIP2	AA203144	8.6	4.4	3.8
57628_at	pelota (Drosophila) homolog	PELO	AA436974	3.6	4.4	1.6
54063_at			AI801777	16.9	4.4	6.2
53810_f_at			AA401302	0	4.4	1.2
79504_at	transmembrane protein 4	TMEM4	AI939329	3.4	4.4	2.5
48355_at			AI025023	0.9	4.4	4.9
48737_at			AI831738	1.3	4.4	6
63026_at	osteoclast stimulating factor 1	OSTF1	AA886976	8.2	4.4	0

Figure 15XX

65114_at			AA90685	0.1	4.4	0
60690_at			AA20328	4.2	4.4	7.2
60503_f_at			AA176985	1.1	4.4	5.7
72348_f_at			AI216132	3.4	4.4	2.7
81793_r_at			AI591416	0.1	4.4	0.5
70323_f_at			AI743616	3.5	4.4	3.4
39672_at	protein tyrosine phosphatase, non-receptor type 7	PTPN7	M64322	3.2	4.4	4.6
48059_at			R61448	5	4.4	3.5
44823_s_at			AA521476	4.7	4.4	4.8
45293_at	hypothetical protein from EUROMAGE 363668		AA631902	5.1	4.4	4
65620_at	hypothetical protein FLJ14005		W52855	4.6	4.4	4.7
31853_at	embryonic ectoderm development	EED	AF080227	3.1	4.3	0
40099_at	rho/rac guanine nucleotide exchange factor (GEF) 2	ARHGEF2	AB014551	8.8	4.3	1.7
40146_at	RAP1B, member of RAS oncogene family	RAP1B	AL080212	9.7	4.3	5.7
31418_at	high-mobility group (nonhistone chromosomal) protein 17-like 1	HMG17L1	Z97055	3.4	4.3	8
1548_s_at	interleukin 10	IL10	U16720	121.9	4.3	3.2
32755_at	actin, alpha 2, smooth muscle, aorta	ACTA2	X13839	3	4.3	6.9
167_at	eukaryotic translation initiation factor 5	EIF5	U49436	17	4.3	0
34782_at	jumonji (mouse) homolog	JMJ	AL021938	4	4.3	1.4
75179_at	ubiquitin-conjugating enzyme E2D 3 (homologous to yeast UBC4/5)	UBE2D3	R81146	4.6	4.3	2.1
89975_r_at	CASP8 associated protein 2	CASP8AP2	AI811300	0.4	4.3	0
77134_at			AA939344	13.5	4.3	1
74867_at			AI038997	3.7	4.3	3
78918_at			AI763172	7.3	4.3	0
79053_at			AI962796	16.4	4.3	1.6
90341_at			H04388	4.9	4.3	0.2
59198_at	hypothetical protein		AF075050	7.2	4.3	3.6
42721_at			AI261490	17.5	4.3	5.9
91350_at			AA989283	20.2	4.3	3.7
54750_at			AA593586	2.2	4.3	9.8
81125_at			AI221390	1.5	4.3	3
83754_at			AI634046	8.8	4.3	1.7
32153_s_at	ubiquitin B	UBB	U49869	8.2	4.3	3.2
86869_at			AA070496	0.6	4.3	1
68163_at			AI076686	2.9	4.3	2.8

Figure 15YY

57613_at	matrix metalloproteinase 25		MMP25	W72721	0	4.3	0
69973_at				AI693345	1.4	4.3	0
69976_at				AI810528	6	4.3	0
63696_at				AA302686	0.3	4.3	2.4
72601_at	G-substrate			AI802033	0.8	4.3	0
44782_at				AA258063	8.2	4.3	5.9
52770_at	LIS1-interacting protein NUDE1, rat homolog			AI857685	3.9	4.3	4.2
89121_at				AI740879	0	4.3	5.3
50228_at	hemoglobin, gamma G		HBG2	AA992592	0	4.3	7.4
73221_at				AA700015	3.7	4.3	0
45721_at	endothelin converting enzyme 1		ECE1	AL039866	12.6	4.3	5.7
61871_r_at	WW Domain-Containing Gene			AI963349	3.5	4.3	1.5
89990_f_at				AW016185	1.2	4.3	0.8
53302_r_at				AI741209	5.6	4.3	10.1
44397_at				H11252	5.1	4.3	4.4
88302_at				AI082708	2.7	4.3	1.9
75192_at	restin (Reed-Steinberg cell-expressed intermediate filament-associated protein)		RSN	AW008377	4.2	4.3	0.2
45929_at				R92031	2.5	4.3	3.4
53588_at				AI967984	29.2	4.3	33.3
47032_at				AI885390	1.5	4.3	13.8
49645_r_at				N66472	4.9	4.3	6.8
57591_r_at	zinc finger protein (ZFD25)			AW003458	3.5	4.3	2.5
45099_at				N92526	6.1	4.3	8.8
62211_at				AI808804	1.3	4.3	11.7
51849_at				AA190571	0.9	4.3	4.5
54020_at	zinc finger protein 161		ZNF161	AA045464	3.7	4.3	8.1
62527_r_at	DAZ associated protein 1		DAZAP1	AA523424	2.3	4.3	1
74823_f_at	histidyl-tRNA synthetase		HARS	AW022495	5.1	4.3	1.8
47107_at				AI765321	3.1	4.3	5.6
55692_at	hypothetical protein FLJ11656			W22924	16.4	4.3	2.7
56072_at				AA939337	0.2	4.3	3.1
52883_at				AI052530	0	4.3	1.7
84596_at				AA813290	1.7	4.3	0.5
38671_at	KIAA0620 protein			AB014520	4.1	4.3	5.3

Figure 15ZZ

51983_at									
53906_at	differentially expressed in FDCP (mouse homolog) 6								
53120_at	hypothetical protein FLJ20442	DEF6							
560_s_at									
32135_at	sterol regulatory element binding transcription factor 1								
32725_at	BH3 interacting domain death agonist	SREBF1							
33227_at	interleukin 10 receptor, beta	BID							
31496_g_at	small inducible cytokine subfamily C, member 2	IL10RB							
41169_at	plasminogen activator, urokinase receptor	SCYC2							
33830_at	leptin receptor gene-related protein	PLAUR							
1070_at	general transcription factor IIB								
37650_at	makorin, ring finger protein, 1	GTF2B							
41562_at	murine leukemia viral (bmi-1) oncogene homolog	MKRN1							
39986_at	DKFZP586D0919 protein	BMI1							
40332_at	7-60 protein								
35830_at	KIAA0370 protein								
37048_at	translocase of outer mitochondrial membrane 34								
890_at	ubiquitin-conjugating enzyme E2A (RAD6 homolog)								
88461_at		UBE2A							
50745_at									
65910_at									
90740_at									
49972_at									
47513_at									
65239_at									
80694_at									
67565_r_at									
63394_at	RBP1-like protein								
43488_at	transporter-like protein								
67103_at									
59008_i_at									
81045_at	zinc finger protein 24 (KOX 17)								
59085_at	hypothetical protein FLJ22609	ZNF24							
68160_at									
43766_g_at									

Figure 15AAA

59126_at			AI660548	10.2	4.2	8
88520_at			T85261	3.1	4.2	2.6
64001_s_at	seven in absentia (Drosophila) homolog 2	SIAH2	AW008929	11.6	4.2	5.3
78478_r_at	FSHD region gene 1	FRG1	H89231	0.6	4.2	0.1
49430_at	g20 protein		AI928807	0	4.2	0
71936_at			AI148006	5.6	4.2	3.2
72553_at			AI335361	14.7	4.2	3.4
72778_at			AI655731	0.2	4.2	3.7
43750_at	tumor necrosis factor receptor superfamily, member 10a	TNFRSF10A	AA102746	10.6	4.2	6.2
70495_at			AI453095	5	4.2	2.8
45479_at	hypothetical protein PRO1580		H26396	7.1	4.2	1.3
47493_at			AI016311	0	4.2	7.4
65556_at	myosin IB	MYO1B	AI041830	3.2	4.2	0.9
51185_at	PAN2 protein		AI309334	5.6	4.2	3.3
45857_at			AI668561	0	4.2	3.5
81091_at			AW043782	54.4	4.2	0.1
90190_at			AI819924	0	4.2	0
69126_at			AA470958	0.6	4.2	4.5
53199_g_at	chromosome 11 open reading frame 14	C11ORF14	AA609076	0.2	4.2	2.3
64128_at			AW043835	0	4.2	1.8
45659_at			AA582196	1.4	4.2	0.3
53010_at			AI809925	28.6	4.2	29.8
46125_at			AW001628	0.9	4.2	2.5
53204_at	hypothetical protein FLJ10808		AA043165	11.5	4.2	0.1
46203_at	chromosome 6 open reading frame 1	C6ORF1	AW026006	1.4	4.2	1
49085_at	vacuolar protein sorting 29 (yeast homolog)	VPS29	AI346485	0.7	4.2	0
54325_at			AI149232	9.2	4.2	0.8
56649_r_at			AA262228	1.2	4.2	2.6
48710_at			AA480092	3.8	4.2	0.1
58827_at			AI992368	3.8	4.2	5.3
48992_at			AA435933	4.4	4.2	6.4
40848_g_at	KIAA0750 gene product		AB018293	4.3	4.2	4
42831_at			AI201843	4.7	4.2	4.7
57328_s_at			AA976857	5.9	4.2	6.9
61735_at			AI433706	2.5	4.2	5.2

Figure 15BBB

47370_at				AA046457	4.4	4.2	4.4
45115_at				AA909042	4.3	4.2	4.5
41235_at	activating transcription factor 4 (tax-responsive enhancer element B67)	ATF4		AL022312	6.7	4.1	3.1
41654_at	adenosine deaminase	ADA		X02994	39.7	4.1	3.2
32142_at	serine/threonine kinase 3 (Ste20, yeast homolog)	STK3		U26424	5	4.1	3.6
151_s_at	tubulin, beta polypeptide	TUBB		V00599	2.5	4.1	0.5
1876_at	v-ral simian leukemia viral oncogene homolog A (ras related)	RALA		M29893	13.9	4.1	2.2
39976_at	hypothetical protein from EUROIMAGE 783647			AL050087	9.4	4.1	5.9
40016_g_at	KIAA0303 protein			AB002301	9.4	4.1	2.5
37718_at	KIAA0096 protein			D43636	2.2	4.1	4.1
38111_at	chondroitin sulfate proteoglycan 2 (versican)	CSPG2		X15998	71.2	4.1	27.4
37900_at	peroxisomal biogenesis factor 11B	PEX11B		AF093670	1.4	4.1	0
38441_s_at	membrane cofactor protein (CD46, trophoblast-lymphocyte cross-reactive antigen)	MCP		X59408	2.9	4.1	4.2
38814_at	ATPase, H+ transporting, lysosomal (vacuolar proton pump), member J	ATP6J		AF038954	1.6	4.1	6.9
39141_at	ATP-binding cassette, sub-family F (GCN20), member 1	ABCF1		AF027302	4.7	4.1	0
63321_at	hypothetical protein FLJ10702			AA207243	5.6	4.1	1
50713_at				H73401	5.8	4.1	6.6
58524_at				W03029	5.5	4.1	0
75804_at				AI378570	1.3	4.1	2.9
77849_r_at				AI499214	0.4	4.1	0.1
43372_f_at	immunoglobulin heavy constant gamma 3 (G3m marker)	IGHG3		N22028	1.6	4.1	2.1
47144_at	ASB-1 protein			AI018173	5.7	4.1	4.5
55194_at				AI638607	7.9	4.1	4.8
81707_at				AI740571	3.6	4.1	2.9
84315_at				AA602942	0.1	4.1	0.8
42350_r_at				N71364	2.3	4.1	8.8
58626_at				AI200948	2.8	4.1	1.5
56889_at	dual specificity phosphatase 10	DUSP10		N36770	14.2	4.1	10.6
58426_at	KIAA0129 gene product			AI761804	2.3	4.1	6.1
48164_at				T72557	5.1	4.1	5.2
75825_at	chromosome 12 open reading frame 6	C12ORF6		AI912757	4.2	4.1	0.2
71013_at	interleukin-1 homolog 1			AI814314	71.5	4.1	2.3

Figure 15CCC

71432_at			AI926493	11.7	4.1	2.4
51074_g_at			N49836	4.6	4.1	0.4
50371_at	KIAA1321 protein		AI743689	6.6	4.1	0
78614_at			AI458853	1.3	4.1	0.2
64189_at	DKFZP564J0863 protein		W60080	13.1	4.1	0
46634_at	triggering receptor expressed on myeloid cells 1 (TREM1)		D78812	3.7	4.1	2.2
52416_at			R69859	1.5	4.1	2.9
70255_at			AI720732	0.4	4.1	4.2
70412_at			AI478179	6.1	4.1	0
51232_at	hypothetical protein FLJ20764		AI684508	62.7	4.1	2
45948_r_at			R98767	4.6	4.1	6.7
54441_at			AI686323	1	4.1	0.4
47830_at	CA11		AI333599	3	4.1	2.4
54610_at			AI479815	23.1	4.1	9.5
46703_at			AI557105	2.3	4.1	2.2
48952_at	hypothetical protein FLJ21634		AI985274	4.9	4.1	0.3
54596_at			AI768162	0.5	4.1	0.3
52912_at			3096753	0	4.1	0.2
36443_at	dynein, axonemal, heavy polypeptide 9	DNAH9	X99947	4.1	4.1	3.7
85498_at			AI470916	4	4.1	3.8
52342_at	GTT1 protein		AI097206	6.1	4.1	9.3
57261_f_at	ribosomal protein, large, P0	RPLP0	AA588862	4.4	4.1	7.9
47640_at	Nedd-4-like ubiquitin-protein ligase		AI668780	3.9	4.1	5.1
37468_at	Janus kinase 2 (a protein tyrosine kinase)	JAK2	AF058925	3.7	4	3.1
39339_at	KIAA0792 gene product		AB018335	1.5	4	0.6
31802_at	KIAA0226 gene product		D86979	0.7	4	1.9
39742_at	TRAF family member-associated NFKB activator	TANK	U59863	5.2	4	3.1
35012_at	myeloid cell nuclear differentiation antigen	MNDA	M81750	0	4	3
1448_at	proteasome (prosome, macropain) subunit, alpha type, 3	PSMA3	D00762	4.3	4	5.6
32820_at	CCR4-NOT transcription complex, subunit 4	CNOT4	U71267	4.7	4	10.4
1478_at	IL2-inducible T-cell kinase	ITK	L10717	9.7	4	7
39219_at	CCAAT/enhancer binding protein (C/EBP), gamma	CEBPG	U20240	6.3	4	1.7
33897_at	hypothetical protein DKFZp761G1923		AF070611	0.5	4	2.9
41349_at	presenilin 2 (Alzheimer disease 4)	PSEN2	L43964	6.5	4	0
36815_at			AF038185	0	4	1.8

Figure 13DDD

38732_at	chloride channel, nucleotide-sensitive, 1A	CLNS1A	X91788	3.6	4	0
56362_at	hypothetical protein FLJ20585		AA121624	10.9	4	0
61865_at			AI423165	7.3	4	3.9
43360_s_at			AA604273	10.9	4	2
90289_at			AI827330	4.6	4	0
73824_at			AI760159	1.6	4	0.7
81825_at			AI143906	12.1	4	0.5
65162_at			T65020	1.9	4	0.1
82357_at			AA135740	2.4	4	1.9
61842_at	hypothetical protein FLJ20716		AI769900	1	4	0.8
55810_at	regulator of differentiation (in S. pombe) 1	ROD1	AA927668	6.1	4	1.8
59249_at			AI218538	2.2	4	0
47898_at			AI566854	0	4	4.2
80829_at			AA210905	5.2	4	3.4
47233_at			AI038387	2.2	4	2.3
91587_at			AI554946	21.1	4	1.9
84715_at			AI307632	10.6	4	1.8
42813_at			AL043172	0.7	4	5.5
62502_at	ELL-RELATED RNA POLYMERASE II, ELONGATION FACTOR		C75510	20.4	4	2.1
84833_at			AI733516	0.4	4	1.2
43225_i_at			AA001052	9.5	4	4.8
72504_at			AA702785	4.1	4	1.9
44838_at	hypothetical protein FLJ12752		AI271348	13	4	3.1
64295_s_at	hypothetical protein FLJ22457		AI983783	1.4	4	0
49243_at			AI341615	7	4	4.2
43967_at	ATP-dependant interferon response protein 1		AI826411	1.2	4	0
72397_f_at			AI570334	1.7	4	1
56309_r_at			T90530	0	4	4.4
87592_r_at			AA581165	2.7	4	4.5
50329_s_at	RAB9, member RAS oncogene family	RAB9	AI934576	9.8	4	2
56366_at			AA700906	3	4	1.5
72591_at			AA693377	2.2	4	2.5
73711_at			AI761952	3.2	4	0.6
64247_at			H40631	20.6	4	3.1
79412_at	hypothetical protein FLJ10903		AL037735	1	4	1.3

Figure 15EEE

75105_f_at	BCL2-associated athanogene	BAG1	AW024313	1.5	4	0.4
45206_at	zinc finger protein 277	ZNF277	AA102582	3.5	4	2.5
45635_at	HT021		AI801898	2.5	4	0.9
64476_at			AL047045	6	4	0
46234_at	CGI-148 protein		AA043242	4.5	4	4.9
47870_r_at	hypothetical protein DKFZp761C169		AW003245	22.9	4	8.8
46042_at			N98235	3.7	4	5.4
43817_f_at	clusterin (complement lysis inhibitor, SP.40,40, sulfated glycoprotein 2, testosterone-repressed prostate message 2, apolipoprotein J)	CLU	AW008412	26.7	4	2
84838_at			H61113	8	4	2.3
57744_at	HSPC072 protein		AI572226	3.2	4	5.8
82532_at			AI821392	50	4	1.7
33857_at	p47		N25122	4	4	3.8
36298_at	peripherin	PRPH	L14565	1.7	4	2.5
33569_at	macrophage lectin 2 (calcium dependent)		D50532	4	4	4
48811_at	N-acetylglucosamine-1-phosphodiester alpha-N-acetylglucosaminidase		AW007594	5.2	4	5
50926_s_at	fatty acid hydroxylase		R54585	4	4	4
44085_at			AL040188	4.4	4	4.6
36495_at	fructose-1,6-bisphosphatase 1	FBP1	U21931	5.4	4	4.9
52841_at			AA731740	4.1	4	4
39425_at	thioredoxin reductase 1	TXNRD1	X91247	1.6	3.9	0.3
612_s_at	2',3'-cyclic nucleotide 3' phosphodiesterase	CNP	M19650	8.3	3.9	5.2
31856_at	glycoprotein A repetitions predominant	GARP	Z24680	5.2	3.9	5.3
41129_at	KIAA0033 protein		D26067	2.1	3.9	0.1
32253_at	arginine-glutamic acid dipeptide (RE) repeats	RERE	AB007927	0	3.9	16.4
33410_at	integrin, alpha 6	ITGA6	S66213	1.3	3.9	12.1
35738_at	high-mobility group (nonhistone chromosomal) protein 17-like 3	HMG17L3	AI347088	6.9	3.9	3.6
39659_at	Ts translation elongation factor, mitochondrial	TSFM	L37936	11.4	3.9	0
36934_at	hypothetical protein LOC57158		AL035447	7.8	3.9	13.6
36942_at	KIAA0174 gene product		D79996	7.3	3.9	1.9
37275_at	GA-binding protein transcription factor, beta subunit 1 (53kD)	GABPB1	U13045	6.5	3.9	10.2
38099_r_at	fatty acid-Coenzyme A ligase, long-chain 4	FACL4	AF030555	8.6	3.9	1.4
37615_at	growth factor receptor-bound protein 10	GRB10	D86962	19	3.9	60.3

Figure 15FFF

38473_at	threonyl-tRNA synthetase	TARS	M63180	7	3.9	1.2
277_at	myeloid cell leukemia sequence 1 (BCL2-related)	MCL1	L08246	5.2	3.9	10
38796_at	complement component 1, q subcomponent, beta polypeptide	C1QB	X03084	4.3	3.9	0
39132_at	SWI/SNF related, matrix associated, actin dependent regulator of chromatin, subfamily a, member 5	SMARCA5	AB010882	7.1	3.9	16.4
33212_at	ribosome binding protein 1 (dog 180kD homolog)	RRBP1	AF006751	2.7	3.9	1.3
51831_s_at			AW001505	1.4	3.9	6.3
48798_at	CTL2 gene		AI569503	2	3.9	1.7
50409_at	KIAA1451 protein		AI674191	3.1	3.9	0
75244_at			AI732578	0.8	3.9	0.4
80657_at			AI392922	12.4	3.9	1
54624_at			AI700571	3.2	3.9	2.3
81052_r_at			AA688189	2	3.9	1.4
65478_at			AI744031	4.5	3.9	1.2
51643_at			AL036254	5.9	3.9	6.5
67903_at			R07848	9.9	3.9	0
42692_at			AA416843	11.1	3.9	2.7
69038_at			AI375077	0.5	3.9	4.2
56279_at	GTF2I repeat domain-containing 1	GTF2IRD1	R12398	0.2	3.9	0
49600_at			AI936984	3.5	3.9	7
75816_at			AA495930	0	3.9	2.5
51035_f_at	hypothetical protein FLJ22672		AA310453	3.4	3.9	3.8
52535_s_at	cyclin-dependent kinase inhibitor 1B (p27, Kip1)	CDKN1B	AI680672	9.1	3.9	8.3
44344_at	H2B histone family, member L	H2BFL	AA557205	26.4	3.9	0.5
44808_at			AA455864	2.5	3.9	7.7
44265_at			H10709	1.9	3.9	7.1
84509_at			AI005100	2.2	3.9	0.3
56500_at	junctional adhesion molecule		AI241578	27.4	3.9	0
78501_at			AI880004	1.2	3.9	0
44213_at			AI559696	2.6	3.9	4.3
58809_at	syntaxin 7	STX7	AW022399	1.7	3.9	0
52164_at	chromosome 11 open reading frame 24	C11ORF24	AA065185	0	3.9	0
44580_at	pregnancy-induced growth inhibitor		AI814233	0	3.9	0
43492_at	hypothetical protein FLJ10439		AI818133	3.7	3.9	0.3
52044_at	hypothetical protein FLJ22649 similar to signal peptidase SPC22/23	FLJ22649	AI801545	1.5	3.9	0

Figure 15GGG

64443_at	EH-domain containing 4	EHD4	AI971203	3	3.9	0
54424_at			AI391585	7.7	3.9	1.3
57237_at	TERF1 (TRF1)-interacting nuclear factor 2	TINF2	W25667	3.1	3.9	1
56023_at			AI041556	6.3	3.9	23.6
46724_at	tensin	TNS	AL046979	5	3.9	0
51444_s_at			2436404	1	3.9	1.8
40409_at	aldehyde dehydrogenase 10 (fatty aldehyde dehydrogenase)	ALDH10	U46689	5.3	3.9	5.1
1361_at	telomeric repeat binding factor (NIMA-interacting) 1	TERF1	U40705	6.2	3.9	2.6
31722_at	ribosomal protein L3	RPL3	AL022326	4.9	3.9	9
1630_s_at	spleen tyrosine kinase	SYK	Z29630	5.1	3.9	4.9
44705_at	hypothetical protein		AA133356	4.7	3.9	4.9
53884_at			AA312905	4.9	3.9	4.7
57497_at			AA419263	4.5	3.9	4.1
52908_at	hypothetical protein FLJ21324		AA865619	4.2	3.9	4.1
38716_at	calcium/calmodulin-dependent protein kinase kinase 2, beta	CAMKK2	AB018330	1.5	3.8	0.1
33499_s_at	immunoglobulin heavy constant gamma 3 (G3m marker)	IGHG3	AF067420	5.9	3.8	17.1
1780_at	Gardner-Rasheed feline sarcoma viral (v-fgr) oncogene homolog	FGR	M19722	2.9	3.8	0.3
32125_at	hypothetical protein LOC57187		AA928996	3	3.8	6.8
32697_at	inositol(myo)-1(or 4)-monophosphatase 1	IMPA1	AF042729	6.6	3.8	1
34128_at	UDP-Gal:betaGlcNAc beta 1,3-galactosyltransferase, polypeptide 5	B3GALT5	AB020337	0	3.8	0.3
38676_at	stress 70 protein chaperone, microsome-associated, 60kD	STCH	AA059408	7.6	3.8	4
39298_at	alpha2,3-sialyltransferase		AB022918	0	3.8	0
36312_at	serine (or cysteine) proteinase inhibitor, clade B (ovalbumin), member 8	SERPINF8	L40377	9.2	3.8	0
34864_at	hypothetical protein		AF070638	0.9	3.8	0
35274_at	adaptor-related protein complex 1, gamma 1 subunit	AP1G1	Y12226	4	3.8	2.5
35739_at	myotubularin related protein 3	MTMR3	AB002369	6.1	3.8	0.5
36176_at	tubulin-specific chaperone c	TBCC	U61234	2.6	3.8	7.4
1295_at	v-rel avian reticuloendotheliosis viral oncogene homolog A (nuclear factor of kappa light polypeptide gene enhancer in B-cells 3 (p65))	RELA	L19067	5.6	3.8	5.3
1453_at	MAD (mothers against decapentaplegic, Drosophila) homolog 2	MADH2	U68018	1.3	3.8	0
38172_at	carbonyl reductase 1	CBR1	AB003151	0	3.8	0
34825_at	TRAF and TNF receptor-associated protein		AL031775	1.1	3.8	2.6
39182_at	epithelial membrane protein 3	EMP3	U87947	4.8	3.8	5
66159_l_at			W56067	2	3.8	5.8

Figure 15HHH

46778_at			AA129377	2.4	3.8	6.9
50787_at			AA527219	3.7	3.8	4.3
65158_at			AI741328	5.4	3.8	1.2
76926_at			AI807691	2.3	3.8	2.4
53777_at	hypothetical protein FLJ22693		AI672353	5.2	3.8	2.2
82428_at			AI610475	3.8	3.8	0
57037_at	kinesin family member 5B	KIF5B	AA057424	0	3.8	0.9
81541_at	natural killer cell receptor 2B4		AI871467	2.3	3.8	7.1
84000_at			AI419726	1.5	3.8	1.3
55253_at			AA621047	4.1	3.8	4.3
42825_at			AA020957	2.2	3.8	6.1
85151_at			AA993736	1.7	3.8	0
49251_f_at	hypothetical protein MGC2628		AI969529	0	3.8	4.6
64309_f_at	amyloid beta (A4) precursor protein (protease nexin-II, Alzheimer disease)	APP	AI625555	2.4	3.8	0.6
84998_at			AI760126	6	3.8	4.6
63279_at			R67641	2.4	3.8	4.3
86606_at			AA884702	0.9	3.8	3.7
69817_at			AI807532	0	3.8	1.4
69879_at			AI688187	0.2	3.8	0
45289_at	novel Ras family protein		AA133248	61.7	3.8	24.3
80391_at	kinesin-like 2	KNL2	AI953141	4.1	3.8	6.4
64568_at	translocase of outer mitochondrial membrane 70 (yeast) homolog A	TOMM70A	D60341	1.9	3.8	3.7
44202_at			AA417797	4.7	3.8	6.1
60823_at			AA191336	2.6	3.8	0.2
89730_at			AI948619	0.5	3.8	1.6
75262_s_at			AW025934	0.2	3.8	0
48413_at	ras GTPase activating protein-like		W46352	0.6	3.8	0
62899_at	cytidine monophosphate-N-acetylneuraminic acid hydroxylase (CMP-N-acetylneuraminic monooxygenase)	CMAH	AI279731	2.8	3.8	3.6
65562_at	twisted gastrulation		AA195009	3.2	3.8	0
88199_at	zinc finger protein 219	ZNF219	AI990459	1.8	3.8	1.7
44763_at			AA280886	9.3	3.8	6.9
44139_at			N31986	2.7	3.8	0.6
61812_at	transcriptional intermediary factor 1 gamma		AA001635	1.7	3.8	2.1

Figure 15III

50320_g_at			AI497833	12	3.8	5.7
60676_at			AI093876	2.5	3.8	5.2
56937_at			AI698117	2.3	3.8	1.3
68652_at			AI431778	38	3.8	0.6
49943_at			AI681558	11.3	3.8	8.3
64054_at	hypothetical protein PP3501		R36266	2.7	3.8	0
68410_r_at			T96375	3.8	3.8	4.9
49140_at			AI244908	61.5	3.8	4.4
53420_at			R94756	1.4	3.8	0
52540_at			AI467915	3.9	3.8	2.6
50978_at	putative secreted ligand homologous to ffx1	hfix	AI338625	117.2	3.8	3.3
44499_at			N67390	7.8	3.8	4.6
62480_at			AI446414	3.1	3.8	12.7
53666_at			AA114900	6.6	3.8	6.4
62524_at			AI951932	2	3.8	4.4
64911_s_at	hypothetical protein FLJ11305		AA854113	29.8	3.8	0
47868_at			AI983739	7.1	3.8	0.6
80610_r_at	follicular lymphoma variant translocation 1	FVT1	AW007121	3.7	3.8	1.8
54907_at			R42787	1.6	3.8	8.2
74571_s_at	carbonic anhydrase III, muscle specific	CA3	AI972237	1.2	3.8	2.5
65960_at			R77323	1.9	3.8	1.6
52355_g_at			AW003924	1.6	3.8	8.1
66674_at			AA931761	2.1	3.8	9.3
71770_at			AI052076	0.3	3.8	2.2
78736_r_at			AI094933	4.1	3.8	5.1
87423_at			C04538	1.7	3.8	0.3
33916_at	imidazoline receptor candidate		AB023192	4.4	3.8	3.6
37002_at	biliverdin reductase B (flavin reductase (NADPH))	BLVRB	D32143	5.2	3.8	5.3
33833_at	spectrin, alpha, non-erythrocytic 1 (alpha-fodrin)	SPTAN1	J05243	5.1	3.8	5.7
61259_at			W31943	3.7	3.8	4
52038_r_at			AI223016	3.4	3.8	3.9
51713_at			AI660536	4.2	3.8	4.2
48094_at			AA531025	4.4	3.8	4.1
51239_at	hypothetical protein FLJ23239		W67828	4.9	3.8	4.4
54878_at			AL048421	4.9	3.8	4.8

Figure 15JJJ

55564_at	spleen tyrosine kinase	SYK	AA521157	3.9	3.8	3.9
34427_g_at	major histocompatibility complex, class I-like sequence	HLALS	U22963	4.9	3.7	0.9
39041_at	dihydrolipoamide S-acetyltransferase (E2 component of pyruvate dehydrogenase complex)	DLAT	Y00978	0	3.7	7.6
31898_at	KIAA0212 gene product	kiaa0212	D86967	5	3.7	2.3
35380_at	TEA domain family member 1 (SV40 transcriptional enhancer factor)	TEAD1	M63896	9.3	3.7	6.9
32079_at	kinesin family member 13B	KIF13B	AB014539	0.3	3.7	0.6
32635_at	transcriptional intermediary factor 1 gamma		AB029036	4.4	3.7	6.1
33219_at	KIAA1097 protein		AB029020	4.4	3.7	1.7
893_at	ubiquitin carrier protein		M91670	2.1	3.7	2.6
41142_at	3-oxoacid CoA transferase	OXCT	U62961	23	3.7	4.5
1792_g_at	cyclin-dependent kinase 2	CDK2	M68520	7.4	3.7	2.9
34246_at			AA418437	7.7	3.7	0.6
843_at	protein tyrosine phosphatase type IVA, member 1	PTP4A1	U48296	13.1	3.7	7.2
33856_at	CAAX box 1	CXX1	Y13374	2.6	3.7	0.4
32160_at	seven in absentia (Drosophila) homolog 1	SIAH1	U76247	8.8	3.7	1.7
33556_at	Huntingtin interacting protein E		AF049611	1	3.7	4.1
34392_s_at	RAB1, member RAS oncogene family	RAB1	AL050268	5.9	3.7	2
39941_at	RAD50 (S. cerevisiae) homolog	RAD50	Z75311	2.7	3.7	1.2
1562_g_at	dual specificity phosphatase 8	DUSP8	U27193	2.9	3.7	7.6
36069_at	KIAA0456 protein		AB007925	0	3.7	4.7
41601_at			AA142964	2.1	3.7	5
35311_at	cellular repressor of E1A-stimulated genes	CREG	AF084523	8	3.7	0.2
37674_at	aminolevulinatase, delta-, synthase 1	ALAS1	Y00451	9.9	3.7	12.5
56351_at			AI436023	2.1	3.7	0
48091_at			AA502943	17.7	3.7	0
66304_r_at			R43631	0.3	3.7	1.5
77215_at			AI733246	1.4	3.7	3.5
74219_at			AA994516	3.1	3.7	2.3
85607_at			AI681283	4.1	3.7	0
77696_at			AI733058	0	3.7	0.3
65877_at	splicing factor 3b, subunit 1, 155kD	SF3B1	AA143047	1.6	3.7	0.7
61489_at			AI718763	12.9	3.7	3.4
61067_r_at			AA470798	4.2	3.7	6.4
90499_at			AI251990	2.9	3.7	0.1

Figure 15KKK

58257_at	KIAA1407 protein		AI686022	0.1	3.7	2
60025_at			AI435572	1.9	3.7	3.3
47525_at	Kreisler (mouse) maf-related leucine zipper homolog	KRML	AI869550	0	3.7	3.5
62314_at			H04394	13.8	3.7	4.6
92089_at			AA984147	1.6	3.7	3.8
41153_f at	catenin (cadherin-associated protein), alpha 1 (102kD)	CTNNA1	AF102803	5.5	3.7	0.1
41222_at	signal transducer and activator of transcription 6, interleukin-4 induced	STAT6	AF067575	2	3.7	0
43105_at			T98171	3.6	3.7	9.2
63074_at			AA570454	4	3.7	4.7
79209_at			N30414	10.3	3.7	0.6
68406_at			W86629	4	3.7	3.7
69169_at			AI990366	2.4	3.7	2
70200_r at			AI735341	3.3	3.7	0.7
45308_at			AA114834	2.5	3.7	4
42893_at			R16983	3.1	3.7	15.7
89556_at			AA909694	0	3.7	1.7
43043_at			AA482037	0	3.7	4
72406_f at			AI352077	2.3	3.7	2.1
51147_at			AA258084	2.6	3.7	2.8
75752_at	hypothetical protein FLJ10033		AI473783	3	3.7	0
56595_at			AA192755	0	3.7	3.2
49769_at	hypothetical protein FLJ13611		R61558	3.8	3.7	1.3
56331_at	hypothetical protein FLJ20559		AA478923	17	3.7	1.4
35820_at	GM2 ganglioside activator protein	GM2A	X62078	0.6	3.7	0
57668_s at			N64025	3.8	3.7	7.2
73574_at			AI290214	4.8	3.7	8.1
60011_at	KIAA1255 protein		H62663	3.4	3.7	0.3
88150_r at			AI201287	2.6	3.7	0.4
54930_at			AI290816	8.1	3.7	2.6
46375_at			AI494113	4	3.7	5.7
82660_at	ninein (GSK3B interacting protein)	NIN	T82318	0	3.7	0.2
45612_at			AI744560	4.9	3.7	5.9
62261_s at	LIM domains containing 1	LIMD1	AI933287	4.5	3.7	1.7
53085_at	serine protease inhibitor, Kazal type, 5	SPINK5	AA534163	0.6	3.7	1.5

Figure 15LLL

48121_at	meningioma expressed antigen 5 (hyaluronidase)	MGEA5	AI344277	6.1	3.7	1.3
58764_at	suppression of tumorigenicity 7	ST7	AI821369	2.2	3.7	3
46660_at	hypothetical protein FLJ10074		AA018399	5.6	3.7	0.1
51949_at			AI982758	2.9	3.7	6.3
55596_at			AA634084	1.9	3.7	1.9
65978_at			AI817130	3.2	3.7	0
71976_at			R62974	0.6	3.7	1.3
79735_at			AA037529	6.8	3.7	3.9
81062_at			AA132765	1.7	3.7	2.8
39416_at	Tax interaction protein 1		U90913	5.3	3.7	5.3
2058_s_at	integrin, beta 5	ITGB5	M35011	4	3.7	3.5
40266_at	KIAA1036 protein		AB028959	3.9	3.7	4.3
1427_g_at	Src-like adapter	SLA	D89077	3.5	3.7	5.1
37010_at	general transcription factor IIA, 2 (12kD subunit)	GTF2A2	AI203737	4.1	3.7	4.3
138_at	mitogen-activated protein kinase kinase kinase 1	MAP4K1	U66464	3.4	3.7	2.8
39135_at	KIAA0767 protein		AB018310	4.5	3.7	4.3
210_at	phospholipase C, beta 2	PLCB2	M95678	4.3	3.7	3.6
48810_at	hypothetical protein DKFZp761H221		AI547243	4.6	3.7	6.5
53848_at			AI422099	4.2	3.7	5.4
65670_at			AA489239	3.6	3.7	3
63793_at	DNA polymerase epsilon p12 subunit		AI797479	5.4	3.7	4.8
62440_at			AI198946	1.6	3.7	3.1
56418_at	uncharacterized hematopoietic stem/progenitor cells protein MDS027		N25128	5.1	3.7	5.8
45339_at	CGI-39 protein; cell death-regulatory protein GRIM19		AA181145	3.3	3.7	4.4
45607_at			AI810669	3.7	3.7	3.6
47390_at			AA928060	3.7	3.7	3.7
38704_at	actin binding protein; macrophin (microfilament and actin filament cross-linker protein)		AB007934	3.8	3.6	4.7
38999_s_at	N-myristoyltransferase 1	NMT1	M86707	1.6	3.6	0.3
41716_at	KIAA0856 protein		AB020663	3.1	3.6	1.6
724_at	adrenergic, beta, receptor kinase 2	ADRBK2	X69117	3.8	3.6	0.5
40200_at	heat shock transcription factor 1	HSF1	M64673	3	3.6	0.6
40235_at	activated p21cdc42Hs kinase		L13738	2.9	3.6	1.8
40470_at	oxoglutarate dehydrogenase (lipoamide)	OGDH	D10523	4.7	3.6	0.3

Figure 15MMM

40803_at				AL050161	6.6	3.6	4.5
37910_at	Rho GTPase activating protein 4	ARHGAP4		U52112	0.7	3.6	1.6
33799_at	seven in absentia (Drosophila) homolog 2	SIAH2		U76248	7.7	3.6	5.4
33803_at	thrombomodulin	THBD		J02973	2.7	3.6	0.2
34689_at	three prime repair exonuclease 1	TREX1		AJ243797	1.9	3.6	1.1
1565_s_at	growth factor receptor-bound protein 2	GRB2		M96995	3.7	3.6	1.1
34948_at				AI820065	1	3.6	3.4
41789_r_at	KIAA0669 gene product			AB014569	3.2	3.6	2.8
41796_at	phospholipase C, epsilon 2	PLCE2		AB029015	1.7	3.6	11.4
40038_at	suppression of tumorigenicity 7	ST7		W02490	0.4	3.6	2.4
36110_at	RAB5A, member RAS oncogene family	RAB5A		M28215	2.4	3.6	3.9
32563_at	ATPase, Na+/K+ transporting, beta 3 polypeptide	ATP1B3		U51478	6.4	3.6	3.8
37553_at	testis-specific kinase 1	TESK1		D50863	1.7	3.6	5
38043_at	glucose-6-phosphate dehydrogenase	G6PD		X55448	1.5	3.6	0
37575_at				AL050192	5.3	3.6	2.1
37921_at	neuronal pentraxin I	NPTX1		U61849	1.3	3.6	10.3
42304_at				AA810277	2.9	3.6	2.1
56940_g_at				AI963304	4.4	3.6	4.3
80994_i_at				T70600	5.7	3.6	0.5
65135_at				AI916853	6.9	3.6	4.9
58872_at				AI983986	0	3.6	0.9
47447_at				AI262829	3.6	3.6	12.9
65166_at				N28835	1.9	3.6	1.1
78001_at				AI139884	1.3	3.6	0
78482_at				AA705165	62.7	3.6	35.7
78751_at				AI097067	2.1	3.6	2.9
80258_at				AW008487	2.2	3.6	2.3
89139_f_at				AI681260	1.7	3.6	2.6
91380_at				AI273546	2.8	3.6	1
91400_at				AI309016	5.4	3.6	3.2
44731_at				AI807804	4.9	3.6	2.2
46744_s_at	carboxypeptidase M	CPM		AA902480	5.3	3.6	3.5
78822_s_at				AI307008	2.8	3.6	0
63406_at	uncharacterized hematopoietic stem/progenitor cells protein MDS030			AI357582	6	3.6	2.4

Figure 15NNN

48597_at			AI871648	1.9	3.6	4.2
70734_at			R14439	0.7	3.6	0.5
63826_at	hypothetical protein FLJ12492		W81045	5.6	3.6	0
73198_at			AL043551	0	3.6	0.5
52226_at			H16779	10.5	3.6	4.2
61273_s_at			AW009229	2.8	3.6	0
50069_at			AI090586	2.3	3.6	5.2
57004_r_at			R67585	2.7	3.6	3.4
42705_at			AI207044	11.2	3.6	2
49895_at			AI061455	2.6	3.6	1.7
44135_at	upstream binding protein 1 (LBP-1a)	UBP1	AI631741	2.3	3.6	0
55341_at			AI351368	3.6	3.6	2.2
49008_g_at			AA399074	1.5	3.6	5.6
52032_at			AA626793	3.1	3.6	1.7
76732_at			N63023	8.8	3.6	1.9
61728_at			AI149793	3.1	3.6	3.2
50971_at			AI523202	0.4	3.6	1.6
45771_at			AI363359	1.9	3.6	0.8
45541_s_at			AW015697	1.7	3.6	7.9
44054_at	KIAA1582 protein		AA489109	9.5	3.6	0.1
52608_at			AI860012	0.3	3.6	0
45580_s_at			AI983200	4.8	3.6	1.8
64692_at			W30810	10.8	3.6	16.2
64909_at	hypothetical protein DKFZp762B226		AI341683	3.3	3.6	1.3
47882_at			H15173	1.7	3.6	7.5
62976_at			AI251931	0.6	3.6	0
34384_at	ATP-binding cassette, sub-family C (CFTR/MRP), member 1	ABCC1	AF022853	1.1	3.6	0
35336_at	KIAA0063 gene product		AL021707	5.3	3.6	6.2
90328_at			AA771960	4	3.6	4
48106_at	hypothetical protein FLJ20489		H14241	4.1	3.6	3.9
81574_at	caspase recruitment domain protein 9		AI307612	4.2	3.6	4.2
38487_at	KIAA0246 protein		D87433	3.8	3.6	3.9
44569_at			AA961420	3.8	3.6	3.6
80251_r_at	ATPase, Ca++ transporting, ubiquitous	ATP2A3	AI380736	3.6	3.6	3.6

Figure 15000

39385_at	alanyl (membrane) aminopeptidase (aminopeptidase N, aminopeptidase M, microsomal aminopeptidase, CD13, p150)	ANPEP	M22324	4.4	3.5	0.1
34740_at	forkhead box O3A	FOXO3A	AF032886	9.9	3.5	6.3
38923_at	FSHD region gene 1	FRG1	L76159	0	3.5	1.1
34521_at	mitogen-activated protein kinase kinase kinase 13	MAP3K13	AB001872	9.3	3.5	1.8
33360_at	f-box and leucine-rich repeat protein 11	FBXL11	AB023221	6	3.5	4.9
39348_at	HMT1 (hnRNP methyltransferase, S. cerevisiae)-like 1	HRMT1L1	X99209	2.4	3.5	5
39267_at	N-acetylglucosamine-phosphate mutase		AF102265	3.7	3.5	7.8
242_at	microtubule-associated protein 4	MAP4	M64571	3.7	3.5	0.4
37617_at	KIAA1128 protein		U90912	5.5	3.5	2.5
41572_r_at	v-rel avian reticuloendotheliosis viral oncogene homolog	REL	X75042	4.8	3.5	3.7
809_at	RAB27A, member RAS oncogene family	RAB27A	U57094	6.3	3.5	5.9
36709_at	integrin, alpha X (antigen CD11C (p150), alpha polypeptide)	ITGAX	Y00093	1.9	3.5	0
40310_at	toll-like receptor 2	TLR2	AF051152	4.7	3.5	0
32628_at	zinc finger protein 161	ZNF161	D28118	3.7	3.5	7.2
35793_at	Ras-GTPase activating protein SH3 domain-binding protein 2		AB014560	1.5	3.5	4.4
40347_at			AA913812	4.9	3.5	0.7
37399_at	aldo-keto reductase family 1, member C3 (3-alpha hydroxysteroid dehydrogenase, type II)	AKR1C3	D17793	3.6	3.5	0.1
37694_at	KIAA0244 protein		D87685	3.8	3.5	0
38391_at	capping protein (actin filament), gelsolin-like	CAPG	M94345	1.3	3.5	0.8
39169_at	Sec61 gamma		AF054184	7.3	3.5	1.2
61737_at			AI750558	2.4	3.5	2.5
66186_at			Z41019	2.8	3.5	0.6
73520_at			AI791312	5.9	3.5	0.9
81655_at			AI824316	0.4	3.5	1.5
74116_at			N71863	0.5	3.5	2.2
79218_at			AA928876	2.4	3.5	0.4
54408_at	zinc finger protein 143 (clone pHZ.1)	ZNF143	AI816009	0.3	3.5	1
42624_at			H19417	1.9	3.5	5.1
81555_at	hypothetical protein FLJ20733		AI360255	0	3.5	0.4
84320_at			AA602573	3.2	3.5	4.8
48405_at			R68394	2.4	3.5	2.6
47948_at			H04453	3.1	3.5	2
62602_at			AA418402	4.2	3.5	0

Figure 15PPP

85186_r_at			AI889871	1.1	3.5	1.5
84410_at	sphingosine-1-phosphate lyase 1	SGPL1	AA036889	5.3	3.5	1.9
46958_at			AI868421	0	3.5	0.9
79562_at			AI829674	3.1	3.5	3.8
65757_at	zinc finger protein 265	ZNF265	AW002312	2.5	3.5	1.8
64345_s_at	KIAA1102 protein		AW003533	3.1	3.5	2.3
70318_i_at			AI741937	0.8	3.5	3.4
89412_at			AI804386	8.3	3.5	5.3
59683_at	neuregulin		AI338338	1.6	3.5	3.4
73928_at			AA505551	4.5	3.5	0.3
50376_at	hypothetical protein FLJ11137		AI278629	2.5	3.5	4.9
55716_at	hypothetical protein FLJ10902		AA194996	3.3	3.5	0.5
44104_at	ERO1 (S. cerevisiae)-like		AL036582	8.1	3.5	0.2
65800_s_at	hypothetical protein		T89723	0.4	3.5	0.8
56850_at			R08417	14.8	3.5	1.2
55023_f_at			H14337	0	3.5	7.9
57739_at	hypothetical protein FLJ20195		AI949010	4	3.5	8.7
56540_at			AI417193	8.4	3.5	0
64120_i_at	hypothetical protein MGC4825		AI459097	7.9	3.5	1.6
72999_f_at	chromosome 8 open reading frame 1	C8ORF1	AW002557	0.9	3.5	0.4
52832_at			AA012862	1.8	3.5	4.3
69914_at			AA447177	6	3.5	2.4
61128_at			AI338787	1.5	3.5	5.4
47028_at			AI732104	3.1	3.5	1.6
45071_at			N69066	3.3	3.5	5.8
57810_at	hypothetical protein		AA825204	11.7	3.5	1.3
54450_at			AI400303	1.9	3.5	3.3
52334_at			3052393	1.6	3.5	2.9
91655_f_at		TGFB1	AI419764	3.1	3.5	18.1
1385_at	transforming growth factor, beta-induced, 68kD	RPS6KA1	M77349	6.4	3.5	6.3
1127_at	ribosomal protein S6 kinase, 90kD, polypeptide 1	HNRPD	L07597	2.6	3.5	3.5
40770_f_at	heterogeneous nuclear ribonucleoprotein D-like	HDAC5	AB017019	4	3.5	3.3
38810_at	histone deacetylase 5		AF039241	4.2	3.5	4.2
53818_at			AI041816	7.3	3.5	4.5
60009_at			AI673085	3.7	3.5	2.6

Figure 15QQQ

88691_at	unr-interacting protein		AI660343	3.1	3.5	3.2
39971_at	lymphoblastic leukemia derived sequence 1	LYL1	M22637	4.8	3.5	4.8
44136_at	hypothetical protein FLJ13910		W68512	4.7	3.5	4.1
54948_at			AI684747	4.1	3.5	3.9
54969_at			AI829724	4.3	3.5	4
37104_at	peroxisome proliferative activated receptor, gamma	PPARG	L40904	2.3	3.4	1.4
39809_at	HMG-box containing protein 1		AF019214	3.8	3.4	2.7
39047_at	KIAA0156 gene product		AB020880	3.7	3.4	2
39860_at	far upstream element (FUSE) binding protein 1	FUBP1	U05040	3.2	3.4	0
32987_at	FSH primary response (LRPR1, rat) homolog 1	FSHPRH1	X97249	3.4	3.4	2
40176_at	ret finger protein	RFP	J03407	6.1	3.4	0
39733_at	homocysteine-inducible, endoplasmic reticulum stress-inducible, ubiquitin-like domain member 1	HERPUD1	AF055001	1.4	3.4	9.1
33598_r_at	chromosome 1 open reading frame 7	C1ORF7	AF054176	9.3	3.4	3.2
38223_at	vascular Rab-GAP/TBC-containing		AB024057	1	3.4	0.5
1681_at	estrogen receptor 1	ESR1	X03635	2.6	3.4	2.5
41134_at	KIAA0964 protein		AB023181	5.3	3.4	1.7
41249_at	guanine nucleotide binding protein (G protein), beta polypeptide 1	GNB1	AL031282	2.8	3.4	2.9
32259_at	enhancer of zeste (Drosophila) homolog 1	EZH1	AB002386	3.4	3.4	3.5
33760_at	peroxisomal biogenesis factor 14	PEX14	AB017546	4.6	3.4	0.2
34178_at	zinc finger protein 297	ZNF297	AI884738	6	3.4	5.3
38703_at	aspartyl aminopeptidase	DNPEP	AF005050	1.1	3.4	0
35328_at			AF055023	6.2	3.4	7.9
36649_at	polymyositis/scleroderma autoantigen 2 (100kD)	PMSCL2	X66113	2.4	3.4	0.6
37004_at	surfactant, pulmonary-associated protein B	SFTPB	J02761	1.8	3.4	1.9
32564_at	protein translocation complex beta		AA083129	9.6	3.4	0.6
37392_at	phosphorylase kinase, beta	PHKB	X84908	0.9	3.4	0
37395_at	ATPase, vacuolar, 14 kD		D49400	0.9	3.4	1.2
37915_at	bromodomain adjacent to zinc finger domain, 2B	BAZ2B	AL080173	8.9	3.4	3.9
55748_at	ribosomal protein L17 isolog		AI141700	1.5	3.4	0.3
57990_g_at			AI206946	0	3.4	0
49935_at			AI886666	0.2	3.4	0.4
48873_at	Gene 33/Mig-6		AI571452	5.2	3.4	25.2
75743_at			AA668789	4.9	3.4	1.6
64903_at			N30209	0	3.4	0.4

Figure 15RRR

51076_at			N49842	0.1	3.4	4.1
88182_at			A1084454	1.9	3.4	1.6
55572_at			AA938267	2.2	3.4	4.1
87633_r_at			AA634506	1.2	3.4	2.2
80194_at			A1859280	3.3	3.4	1.3
65255_r_at			AA393998	3.8	3.4	2.5
68250_at			AA668871	10.6	3.4	1.9
82450_at	T-box 21		A1762808	2.2	3.4	1
65935_at	translocase of inner mitochondrial membrane 8 (yeast) homolog B	TIMM8B	AW005802	0.3	3.4	2.3
81092_g_at	aminopeptidase		R22164	0.6	3.4	1.1
47452_r_at			R39206	0.4	3.4	2.2
44148_at			AA745720	1.6	3.4	2.1
78368_at			A1022066	2.7	3.4	1.5
62209_at			AA860391	1.9	3.4	1.8
46444_at			A1288239	0.7	3.4	4.2
65887_at	transducer of ERBB2, 2	TOB2	A1830948	4	3.4	5
45482_s_at			N93263	4.7	3.4	0
60638_at			W74633	2.7	3.4	3.6
55292_at			W95037	1.4	3.4	2.5
58617_at			AA149312	5.1	3.4	0.3
51902_at	hypothetical protein		AA215376	0	3.4	0
57660_r_at			R43504	4.1	3.4	5.6
55924_at			AA085776	1.3	3.4	0
56523_at			AA194120	2.7	3.4	1.1
62572_at			A1140917	0	3.4	3.7
43417_r_at	CBF1 interacting corepressor		AA199649	2.8	3.4	3.4
55588_at	hypothetical protein FLJ13868		A1814485	17.1	3.4	1.7
63872_g_at	hypothetical protein FLJ10948		AW025350	1.5	3.4	2.6
45722_at			A1547173	1.2	3.4	0
89004_at			A1735078	3.5	3.4	1.8
35343_at	glutamic-oxaloacetic transaminase 1, soluble (aspartate aminotransferase 1)	GOT1	M37400	3.5	3.4	2.9
1357_at	ubiquitin specific protease 4 (proto-oncogene)	USP4	U20657	5.7	3.4	8.3
52238_s_at	transforming, acidic coiled-coil containing protein 3	TACC3	A1990642	3.9	3.4	3.5
56238_at	PDZ-LIM protein mystique		A1589776	3.5	3.4	3.5

Figure 15TTT

49005_at	myosin regulatory light chain interacting protein		AI954822	3.8	3.4	3.6
74047_i_at			AW023188	2.9	3.4	2.5
57327_s_at	ribonuclease, RNase A family, k6	RNASE6	AW026402	4	3.4	3.7
45285_at	proteasome (prosome, macropain) subunit, alpha type, 7	PSMA7	AA877820	4.2	3.4	5.1
47982_at	aquaporin 3	AQP3	AA630981	3.5	3.4	3.1
60873_at	3'-phosphoadenosine 5'-phosphosulfate synthase 2	PAPSS2	AA643513	1	3.4	3.7
45543_r_at	mitochondrial import receptor Tom22		AI359916	3.4	3.4	2.8
54483_at			AA760777	3.4	3.4	3.4
55611_at	hypothetical protein FLJ20011		AA242757	3	3.4	3.4
38659_at	suppressor of clear, C. elegans, homolog of		AB020669	3.5	3.3	4.8
38688_at	KIAA0461 protein		AB007930	3.1	3.3	7.4
41432_at	ribosomal protein S6 kinase, 90kD, polypeptide 5	RPS6KA5	AF090421	5.5	3.3	8.7
39378_at	beclin 1 (coiled-coil, myosin-like BCL2-interacting protein)	BECN1	U17999	5.7	3.3	0.1
32130_at	actin, alpha 1, skeletal muscle	ACTA1	W25984	1.7	3.3	2.7
32961_at	c-myc promoter-binding protein		X63417	6.3	3.3	2.1
39200_s_at			W28760	0	3.3	2.3
38143_at	kallikrein 7 (chymotryptic, stratum corneum)	KLK7	L33404	1.2	3.3	0
40462_at	DKFZP727M231 protein		AF055022	3.8	3.3	0
34672_at	KIAA0603 gene product		AB011175	2	3.3	1.6
33891_at	chloride intracellular channel 4	CLIC4	AL080061	9	3.3	1.8
34671_at	polymerase (RNA) III (DNA directed) (62kD)		U93867	2.2	3.3	0.8
1496_at	protein tyrosine phosphatase, receptor type, A	PTPRA	M34668	2.5	3.3	0.4
35215_at	HDCMA18P protein		AL049996	2.2	3.3	2.6
35305_at	X-prolyl aminopeptidase (aminopeptidase P)-like	XPNPEPL	X95762	2.2	3.3	0
688_at	proteasome (prosome, macropain) 26S subunit, ATPase, 1	PSMC1	L02426	0.9	3.3	1.3
41597_s_at	SEC22, vesicle trafficking protein (S. cerevisiae)-like 1	SEC22L1	AF047442	7.4	3.3	0
36122_at	proteasome (prosome, macropain) subunit, alpha type, 6	PSMA6	X59417	7.7	3.3	4.7
409_at	tyrosine 3-monooxygenase/tryptophan 5-monooxygenase activation protein, theta polypeptide	YWHAQ	X56468	5.5	3.3	6.2
37737_at	protein-L-isoaspartate (D-aspartate) O-methyltransferase	PCMT1	D25547	0	3.3	0
40704_at	phosphoinositide-3-kinase, catalytic, alpha polypeptide	PIK3CA	Z29090	2.9	3.3	1.3
89729_at			AA216357	2.9	3.3	0
48088_at			AI300073	0	3.3	4.7
61757_at			AI273692	8	3.3	7.4
63364_at			AI798098	2.2	3.3	9.5

Figure 15UUU

76140_at			AI688663	3.6	3.3	2.9
66511_at			AA815264	0.8	3.3	3.3
66774_at			AA978204	5.5	3.3	2.1
74772_at			AI424825	0.4	3.3	0.5
74947_at			AA702963	1.9	3.3	0.8
78695_at			AW001771	0	3.3	0
50602_at			AI221136	1.6	3.3	1.8
79310_r_at			AI869278	1.9	3.3	0.7
89164_at			AI393930	1	3.3	0.5
90033_at	phosphodiesterase 2A, cGMP-stimulated	PDE2A	T66157	0.6	3.3	0.1
53912_at	sorting nexin 11		AI668643	2.6	3.3	0.1
54658_at	hypothetical protein FLJ10955		AI961496	4.9	3.3	1.5
59422_at			AI478332	1.7	3.3	1.2
52729_at	CD44 antigen (homing function and Indian blood group system)	CD44	AA037772	35.3	3.3	0.6
85907_at			AI276085	0.8	3.3	1.1
83378_at			T54678	8.3	3.3	6.3
67046_at			AI628081	2.9	3.3	0.6
189_s_at	plasminogen activator, urokinase receptor	PLAUR	U09937	18.8	3.3	5.3
63986_at	death-associated protein 6	DAXX	AW002820	1.3	3.3	1.5
43697_at			AA504249	7.2	3.3	3.4
49234_s_at	KIAA1389 protein		AI436735	0.4	3.3	4.3
60513_at			AW007066	0	3.3	2.7
63108_at			AI741779	3.8	3.3	2.2
85826_at			AA584865	2.7	3.3	2.7
69698_at			H39848	0	3.3	1.5
69404_at			AI692201	1.3	3.3	0.8
70223_at			AL037324	2.3	3.3	4.7
49454_at	hypothetical protein FLJ10036		AA824298	0.8	3.3	2
69053_at	transcription factor BMAL2		AI688570	17.8	3.3	0.8
72721_at			AI830607	47.1	3.3	1.4
61130_at			AI674404	20	3.3	1.9
65484_f.at	glutaminase	GLS	AI768354	4.3	3.3	0.4
52428_at	regulatory solute carrier protein, family 1, member 1	RSC1A1	AI268381	2.8	3.3	1.9
44064_g.at	TIA1 cytotoxic granule-associated RNA-binding protein	TIA1	AA639796	1.6	3.3	2.7
45328_at	G protein-coupled receptor kinase 7	GPRK7	AI922109	1.4	3.3	3.2

Figure 15VVV

81677_at									
51026_at	hypothetical protein PRO0813								
87107_at									
66109_f_at	hypothetical protein DKFZp547H236								
47508_at	centaurin-alpha 2 protein								
89494_r_at									
72742_at									
64015_g_at									
45919_at									
51058_at									
55809_at									
74930_at	DnaJ (Hsp40) homolog, subfamily B, member 2								
56017_at		DNAJB2							
55081_at	hypothetical protein bA395L14.2								
47128_at									
88448_at	hypothetical protein FLJ11113								
55662_at									
52641_at	putative G protein-coupled receptor								
43069_at	nuclear factor of kappa light polypeptide gene enhancer in B-cells inhibitor, beta	NFKBIB							
1089_i_at									
1339_s_at									
41865_at									
35331_at	catenin (cadherin-associated protein), alpha-like 1	CTNNAL1							
38041_at	UDP-N-acetyl-alpha-D-galactosamine:polypeptide N-acetyltransferase 1 (GalNAc-T1)	GALNT1							
33135_at	solute carrier family 19 (folate transporter), member 1	SLC19A1							
80436_at									
62850_at									
67162_r_at									
52081_at	ZAP3 protein								
51906_at									
44047_at	hypothetical protein FLJ20160								
51056_at	hypothetical protein DKFZp434D0412								
43992_at									

Figure 15WW

44005_at				Al670935	3.3	3.3	3
49326_at	matrix Gla protein	MGP		Al140620	3.5	3.3	3.7
53212_at				R55295	3.3	3.3	2.6
53159_at	hypothetical protein FLJ10829			Al056890	3.1	3.3	3.2
57210_at				Al862775	3.9	3.3	4
46247_at	succinate-CoA ligase, GDP-forming, alpha subunit	SUCLG1		AA826261	4.1	3.3	5.5
55446_at				Al199032	3.5	3.3	3.4
87754_g_at	serologically defined colon cancer antigen 3	SDCCAG3		W02630	3.6	3.3	3.6
65299_at				C15819	3.1	3.3	3.1
65862_at				Al525160	3.7	3.3	3
41099_at	ligase III, DNA, ATP-dependent	LIG3		X84740	2	3.2	5.1
37424_at	HCR (a-helix coiled-coil rod homologue)			AB029343	5.8	3.2	3.5
34470_at	transcription factor EC	TFEC		D43945	3.4	3.2	0
32477_at	calpain 9 (NCL-4)	CAPN9		AF022799	0.2	3.2	0.7
39084_at	enolase 3, (beta, muscle)	ENO3		X56832	1.2	3.2	8.4
41690_at				AL049471	10.7	3.2	5.4
41709_at	PTPRF interacting protein, binding protein 2 (liprin beta 2)	PPFIBP2		AF034803	1.5	3.2	0
34951_at	putative chemokine receptor; GTP-binding protein			D10923	6.3	3.2	4.6
1499_at	farnesyltransferase, CAAX box, alpha	FNTA		L10413	0	3.2	0.7
40097_at	eukaryotic translation initiation factor 1A, Y chromosome	EIF1AY		AF000987	1.6	3.2	10.5
40108_at	KIAA0005 gene product			D13630	5.5	3.2	0.9
32294_g_at	luteinizing hormone/choriogonadotropin receptor	LHCGR		M63108	3.9	3.2	7.5
41225_at	dual specificity phosphatase 3 (vaccinia virus phosphatase VH1-related)	DUSP3		AL049417	0.1	3.2	0
32784_at	serine/threonine-protein kinase PRP4 homolog			AB011108	6.6	3.2	3.3
34196_at				Al337901	0.2	3.2	0
33869_at				AL080218	5.2	3.2	3.8
41335_at	DKFZP566O1646 protein			AL050084	2	3.2	0
32544_s_at	Ras suppressor protein 1	RSU1		L12535	0.6	3.2	0
37367_at	ATPase, H+ transporting, lysosomal (vacuolar proton pump) 31kD	ATP6E		X76228	2	3.2	0
38020_at	KIAA0652 gene product			AB014552	3.5	3.2	1.8
32792_at	GCIP-interacting protein p29			AL031432	5.1	3.2	4.1
38379_at	glycoprotein (transmembrane) nmb	GNMB		X76534	1.3	3.2	1
40686_at	neuromedin B	NMB		Al985272	2.9	3.2	16.9
37922_at	transcobalamin II; macrocytic anemia	TCN2		L02648	0.9	3.2	3.8

Figure 15XXX

37932_at	thyroid hormone receptor interactor 11	TRIP11	Y12490	1.8	3.2	0.4
39150_at	ring finger protein 11	RNF11	U69559	3.9	3.2	6.1
48026_at			AI984814	5.9	3.2	1
58131_at			AI806586	6.3	3.2	3.9
56851_at			AL046179	1.3	3.2	1.8
56959_at			AI376649	2.9	3.2	3.7
76243_at			AA744518	2.7	3.2	2
64704_at			AI378126	2.4	3.2	3.1
76912_at			AI081271	0	3.2	0.1
74170_at			AI793316	3.2	3.2	2
66893_f_at			AA904929	1.9	3.2	3.3
57688_at			AA776462	1.1	3.2	0
42720_at			AI393727	9.5	3.2	11.3
65440_at	SCO (cytochrome oxidase deficient, yeast) homolog 1	SCO1	AA205413	0.9	3.2	0
83201_at			AI984360	1.4	3.2	7.7
47911_at			AI249173	0	3.2	0
82493_at			AI792453	2.8	3.2	1.3
40767_at	tissue factor pathway inhibitor (lipoprotein-associated coagulation inhibitor)	TFPI	M59499	15.5	3.2	1.2
85365_at			AI493870	0	3.2	1.7
49312_at	CGI-102 protein		AA747303	4.4	3.2	7.4
51674_at			N28262	3.8	3.2	0.9
59082_at			AI802955	1.8	3.2	0
67890_at			AI613399	4.4	3.2	1.3
85526_at			R64696	4.8	3.2	0
49383_at	CGI-149 protein		AA167070	3.1	3.2	0.7
79309_at			AA703525	3.1	3.2	0.7
43630_at			AA648993	1.3	3.2	2.2
68708_at			AI560174	0.8	3.2	0.5
49903_at	KIAA1404 protein		W72798	7.5	3.2	6.5
59640_at	zinc finger protein, subfamily 2A (FYVE domain containing), 1	ZNFN2A1	N81203	1.6	3.2	2.5
60791_at	hypothetical protein FLJ11011		AI828160	0	3.2	0
42803_at	E2F transcription factor 5, p130-binding	E2F5	AI377305	13.2	3.2	2.2
61277_at	peptide transporter 3		AI141684	0.7	3.2	0
43489_at	hairy/enhancer-of-split related with YRPW motif-like	HEYL	AL040197	4.4	3.2	5.1

Figure 15YYY

42938_at				R85261	2.5	3.2	3.9
85971_at				N54973	1.7	3.2	5.4
65565_at	GTP-binding protein Sara			A1346486	2.3	3.2	0
48266_at	hypothetical protein FLJ10493			AA196635	8.9	3.2	1.2
49624_at				A1768630	1.1	3.2	0
63361_at	spodin 2, extracellular matrix protein		SPON2	AA570505	3.5	3.2	3.2
65351_r_at				AA146979	5.9	3.2	5.1
61901_at				A1139993	2.7	3.2	0.8
64124_at				AA206408	13.9	3.2	0.5
64101_at				N25092	2.1	3.2	1.9
64107_at	x 001 protein			A1654625	0.4	3.2	1.2
65092_at	hypothetical protein FLJ22555			A1078595	2.8	3.2	1.3
64111_at	WAP four-disulfide core domain 1		WFDC1	A1870778	1.5	3.2	0.7
58564_s_at				N78066	3.4	3.2	3.5
45165_at				A1831506	7.8	3.2	4.8
45216_at				N21420	1.1	3.2	1
44866_s_at	ubiquilin 2		UBQLN2	A1190896	9.8	3.2	0.7
57824_r_at				A1973077	1.6	3.2	5.1
49634_at				N51441	0	3.2	14.9
62165_at	x 006 protein			W72231	4.8	3.2	5.9
64468_at	hypothetical protein FLJ10305			A1075995	2.6	3.2	0
46198_at				A1356548	2.5	3.2	7.9
46629_at	hypothetical protein FLJ12154			A1431797	18.1	3.2	2.2
86353_at				A1767724	0.6	3.2	2.5
55384_g_at				AA454036	3	3.2	4.2
45321_at	Homer, neuronal immediate early gene, 1B			N25931	1.4	3.2	0.5
47618_at				AA600853	1.2	3.2	1.5
53029_at				3145376	2.3	3.2	0.9
85103_at				A1674959	0	3.2	1.1
40504_at	paraoxonase 2		PON2	AF001601	3.5	3.2	2.5
250_at	nuclear factor of activated T-cells, cytoplasmic, calcineurin-dependent 3		NFATC3	L41067	3.3	3.2	3.3
33920_at	diaphanous (Drosophila, homolog) 1		DIAPH1	AF051782	4.3	3.2	3.8
41504_s_at	v-maf musculoaponeurotic fibrosarcoma (avian) oncogene homolog		MAF	AF055376	3.9	3.2	4.1

Figure 15ZZZ

34811_at	ATP synthase, H+ transporting, mitochondrial F0 complex, subunit c (subunit 9) isoform 3	ATP5G3	U09813	5.2	3.2	6.2
38107_at	unc119 (C.elegans) homolog	UNC119	U40998	2.9	3.2	2.2
48084_at			AA582193	3.7	3.2	3.7
48833_at	HSPC040 protein		AI749098	4.1	3.2	4.1
59595_at			AA045175	3.6	3.2	3.2
43053_g_at			AW025012	4.9	3.2	4.6
57156_at	HSPC023 protein		W92964	3.4	3.2	4.9
46623_at			AA579641	3.3	3.2	3.3
45224_at			AA025862	5.1	3.2	3.5
57461_at			AI472111	3.5	3.2	3.4
45531_at			AI970101	3.5	3.2	2.5
56267_at	adaptor-related protein complex 1, sigma 2 subunit	AP1S2	AA151802	3.9	3.2	3.7
55678_at	hypothetical protein MGC10986		W72885	3.6	3.2	3.7
65923_at			AI344311	3.7	3.2	4.2
49197_at	solute carrier family 16 (monocarboxylic acid transporters), member 1	SLC16A1	AI129459	3.8	3.2	2.7
54935_at	CD74 antigen (invariant polypeptide of major histocompatibility complex, class II antigen-associated)	CD74	N22918	4	3.2	3.7
37842_at	l-mfa domain-containing protein		AF054589	1.5	3.2	4.7
38666_at	pleckstrin homology, Sec7 and coiled/coiled domains 1 (cytohesin 1)	PSCD1	M85169	6.7	3.1	2.2
39030_at	Rab acceptor 1 (prenylated)	RABAC1	AJ133534	1.6	3.1	3.4
31742_at	tumor necrosis factor (ligand) superfamily, member 14	TNFSF14	AF064090	0.6	3.1	0.1
31845_at	E74-like factor 4 (ets domain transcription factor)	ELF4	U32645	3.5	3.1	1.1
39778_at	mannosyl (alpha-1,3-)-glycoprotein beta-1,2-N-acetylglucosaminyltransferase	MGAT1	M55621	0.4	3.1	0.5
39800_s_at	HS1 binding protein		U68566	7.2	3.1	1.5
31899_at	KIAA0103 gene product		D14659	1.2	3.1	0
32033_at	choline kinase-like	CHKL	AL096780	1.7	3.1	0
40791_at	polymerase (RNA) II (DNA directed) polypeptide A (220kD)	POLR2A	X63564	5.6	3.1	4.2
40952_at	BANP homolog, SMAR1 homolog		AA257983	1.8	3.1	5.7
33354_at	E3 ubiquitin ligase SMURF2		AA630312	1.1	3.1	1.2
34707_at	chromodomain helicase DNA binding protein 3	CHD3	U91543	1.4	3.1	2.7
39576_at	phosphodiesterase 6B, cGMP-specific, rod, beta (congenital stationary night blindness 3, autosomal dominant)	PDE6B	S41458	0.8	3.1	7.7

Figure 15A AAA

87791_at			AA670223	0	3.1	0
88247_at			AI075772	2.7	3.1	1.9
69247_at			AI984074	2	3.1	0.8
69943_at			AI698731	5.2	3.1	2.1
69476_at			AI707474	1.4	3.1	1.7
56263_at			N92517	2.6	3.1	2.5
34204_at	sema domain, immunoglobulin domain (lg), short basic domain, secreted, (semaphorin) 3E	SEMA3E	AL050043	0.4	3.1	4.2
71144_at	tumor necrosis factor receptor superfamily, member 18	TNFRSF18	AI923712	7.7	3.1	0.5
42875_r_at			T97225	2.3	3.1	5.9
45315_at	hypothetical protein FLJ21952		AI825896	0.2	3.1	1.3
44244_at	hypothetical protein FLJ11336		R51067	9.8	3.1	1
65546_at			N78362	0	3.1	5.6
47003_s_at			AW014800	9.3	3.1	0
46587_at			AI142857	2.1	3.1	3.1
43865_at			R49396	1	3.1	5.7
74870_at	carnitine palmitoyltransferase I, liver	CPT1A	AI675101	0.3	3.1	0.4
45719_at			AA541736	2.8	3.1	0
44115_at	hypothetical protein from EUROIMAGE 588495		AA486504	27.1	3.1	3.8
60527_at	Rho GTPase activating protein 5	ARHGAP5	AI768264	3.2	3.1	5.3
54622_at	receptor-interacting serine-threonine kinase 3	RIPK3	N99896	0.9	3.1	0
50976_at			AI817292	12	3.1	0
44588_at			AA604268	4.7	3.1	2.4
44631_at			AI653257	0.2	3.1	3.7
55548_at			AI884542	2.9	3.1	2.2
55492_at			AI858497	1.8	3.1	1.6
45744_at	PRO1912 protein		AL047839	1	3.1	1.2
43952_at			AI721219	0.3	3.1	0
53304_at			R82296	1	3.1	1.8
65477_at	BCL2-like 1	BCL2L1	T15735	12.4	3.1	1.5
61712_at	hypothetical protein PRO0823		AI990512	0.4	3.1	0
51018_at			AA707221	2.5	3.1	1
49054_at	adenylate kinase 3 alpha like		N32441	5.1	3.1	0.4
45598_at	heparanase	HPSE	N45367	1.2	3.1	3.7
66042_at			W87429	0	3.1	1.3

Figure 15CCCC

45620_at			AI623305	2.6	3.1	2.8
45810_at			AI750909	0	3.1	3.1
51857_at			AA705016	0.6	3.1	0
46187_at			AI344122	2.1	3.1	28.9
54487_at			AA044626	3	3.1	5.1
46596_at			AI887633	2.4	3.1	0.4
46608_r_at			AI885018	3.5	3.1	0
64918_at			AI684951	1.2	3.1	2.1
57217_at			AA526961	0	3.1	0
47589_at	similar to acetyl-coenzyme A synthetase	BPHL				
42362_at	biphenyl hydrolase-like (serine hydrolase; breast epithelial mucin-associated antigen)	FGB	AI057008	3.6	3.1	7.1
49743_at	fibrinogen, B beta polypeptide	CEBPD	AI633245	13.4	3.1	3.4
54368_at	CCAAT/enhancer binding protein (C/EBP), delta		AA215655	2.3	3.1	4.2
61398_at			T83269	0	3.1	1.2
47641_at			AA402106	2.5	3.1	2.5
51434_at	KIAA1484 protein		AI732810	1	3.1	4.6
74329_at			2849405	2.1	3.1	2.2
90697_r_at			H92158	3.6	3.1	1.5
41103_at	bromodomain and PHD finger containing, 1	BRPF1	AI345703	3.2	3.1	1.2
38700_at	cysteine and glycine-rich protein 1	CSRPI	M91585	4.1	3.1	3.7
38976_at	coronin, actin-binding protein, 1A	CORO1A	M33146	3.4	3.1	4.4
39774_at	oxidase (cytochrome c) assembly 1-like	OXA1L	D44497	3.1	3.1	3
38956_at	N-acetylglucosamine-1-phosphodiester alpha-N-acetylglucosaminidase		X80695	3.2	3.1	5.4
32317_s_at	sulfotransferase family, cytosolic, 1A, phenol-preferring, member 2	SULT1A2	AF052111	4.1	3.1	3.9
35723_at	nudix (nucleoside diphosphate linked moiety X)-type motif 1	NUDT1	U34804	4.5	3.1	4.1
40024_at	src homology three (SH3) and cysteine rich domain	STAC	D16581	4	3.1	3
35338_at	paired basic amino acid cleaving enzyme (furin, membrane associated receptor protein)	PACE	D86640	3	3.1	3.3
41050_at	TATA box binding protein (TBP)-associated factor, RNA polymerase II, D, 100kD	TAF2D	X17094	2.8	3.1	3.5
58412_at	retinal degeneration B beta		X95525	4	3.1	2.6
47463_at	eukaryotic translation initiation factor 2C, 1	EIF2C1	AA807607	1.4	3.1	3.4
57030_at			AA453723	3.2	3.1	2.9
			W26295	3.1	3.1	3.1

Figure 15DDDD

57823_at				H18472	2.7	3.1	1.8
62889_at	KIAA0675 gene product			D29070	3.1	3.1	3.1
49463_at	suppressor of fused			AI760042	3.3	3.1	3
44343_s_at				AI074877	3.9	3.1	3.8
44143_at	McKusick-Kaufman syndrome		MKKS	AA399076	2.6	3.1	3.5
53082_at				AI741458	4.1	3.1	5.7
53940_at				AI953658	3.3	3.1	3.4
57203_at	hypothetical protein FLJ10913			AA142864	2.5	3.1	3.6
65961_at	CGI-135 protein			W63805	3.1	3.1	3.5
39519_at	KIAA0692 protein			AB014592	7	3	0
32402_s_at	symplesin; Huntingtin interacting protein I			Y10931	2.9	3	0
37435_s_at	potassium inwardly-rectifying channel, subfamily J, member 6		KCNJ6	U52153	5.3	3	6.8
41710_at	hypothetical protein			AL079277	0.7	3	0
925_at	interferon, gamma-inducible protein 30		IFI30	J03909	1.1	3	1.7
33234_at	KIAA0117 protein			AA887480	2.7	3	3.9
1399_at	transcription elongation factor B (SIII), polypeptide 1 (15kD, elongin C)		TCEB1	L34587	0.3	3	0
40956_at	Conserved gene telomeric to alpha globin cluster			X90857	0.7	3	2.1
38944_at	MAD (mothers against decapentaplegic, Drosophila) homolog 3		MADH3	U68019	0	3	0.4
1959_at	antizyme inhibitor			D88674	2.5	3	3.7
34718_at	insulin-like growth factor 1 receptor		IGF1R	X04434	18	3	7.4
34768_at				AL080080	1.9	3	0.5
41489_at	transducin-like enhancer of split 1, homolog of Drosophila E(sp1)		TLE1	M99435	2.1	3	2.7
36005_at	suppressor of white apricot homolog 2			AF042800	1.8	3	2.7
34844_at	KIAA0222 gene product			D86975	1.1	3	2.4
35265_at	fragile X mental retardation, autosomal homolog 2		FXR2	U31501	3.7	3	0.4
36451_at				AI743299	10.6	3	2.3
35348_at	protein kinase, AMP-activated, beta 1 non-catalytic subunit		PRKAB1	AF022116	10.8	3	5.9
37308_at	KIAA1624 protein			AI888084	2.5	3	4.7
35847_at	ubiquitin specific protease 24		USP24	AB028980	3.7	3	4.7
37015_at	aldehyde dehydrogenase 1, soluble		ALDH1	K03000	1.8	3	5.2
1052_s_at	CCAAT/enhancer binding protein (C/EBP), delta		CEBPD	M83667	15.2	3	7
37374_at	annexin A4		ANXA4	M82809	0	3	3.7
38082_at	KIAA0650 protein			AB014550	3.7	3	12.1
38354_at	CCAAT/enhancer binding protein (C/EBP), beta		CEBPB	X52560	3.3	3	0.4

Figure 15EEEE

46750_s_at	hypothetical protein PRO2219		AA479496	2.1	3	3.6
56371_at			AA534298	5.2	3	0
56919_at	KIAA1449 protein	KIAA1449	AI806628	4.2	3	2.2
81429_at	ethanolamine kinase		AI003792	3.7	3	0
76283_at			AA618303	8.5	3	1.1
90346_at			AW003138	0	3	0
58154_at			R38652	2.3	3	0.6
58893_at			AI697856	3.7	3	2.5
76909_at			AI243872	4.1	3	0
66880_at			AI733050	3	3	5.7
78715_at			AI379702	2.3	3	4.8
79343_at			AA525367	2.9	3	3
59210_at			T79178	2.7	3	4.8
52152_at			AI150530	2.9	3	1.5
79970_at			AI733828	1.4	3	2
46457_at	splicing factor (CC1.3)		AI400626	4.6	3	1.3
80539_at			AI700768	1.1	3	0.5
80623_s_at			AI640222	3.1	3	2.4
47108_at	LIM protein (similar to rat protein kinase C-binding enigma)		AA196325	0	3	4.2
81217_at			AI971594	0.6	3	1.1
55191_at	PRO1848 protein		AI334614	2.3	3	2.6
59344_at			AA017033	6.9	3	4.8
84352_s_at			AA971429	2.5	3	1.1
85194_at			AI378091	1.5	3	3.2
44632_at			AA007697	0	3	0.1
59612_at	BH3 interacting domain death agonist	BID	AI669308	8.8	3	0
92136_r_at			AI915601	0.8	3	4.8
75317_s_at	interferon gamma receptor 2 (interferon gamma transducer 1)	IFNGR2	AI985833	0.9	3	0
59775_at			R07483	7.3	3	2.2
68030_at			AI080360	2.4	3	0.6
54863_at			AI215926	0	3	2
68782_f_at			AA635181	2.2	3	0.6
68538_at			AI973024	1.6	3	6.1
53066_at			AI824034	2.8	3	2.9
42475_at			T94540	2	3	4

Figure 15FFFF

70682_at	YME1 (S.cerevisiae)-like 1	YME1L1	AI832869	0.4	3	15.1
72481_at			AI640149	2.8	3	2.4
43032_at	tropomodulin 3 (ubiquitous)	TMOD3	AI373630	3.3	3	0
66103_at			AW008015	6.8	3	2.4
49034_at	CGI-69 protein		AW008524	2.8	3	3.9
53888_at	cathepsin B	CTSB	R81090	0	3	0
45537_at	aquaporin 7	AQP7	F27435	2.2	3	3.1
51096_at	ERO1 (S. cerevisiae)-like		AL048651	8.2	3	0.2
42904_at			R85291	4.7	3	2.3
43990_at	uncharacterized bone marrow protein BM040		AI676178	3.7	3	0.8
51149_at			AA421991	9.7	3	0
50694_at	Carbonic anhydrase-related protein 10		AI815895	4.2	3	4.6
88143_at			AI253134	3.1	3	4.1
54491_at			AA043562	27.5	3	3.1
59668_at			AI066756	2.9	3	0.5
61360_at			AI700588	2	3	2.7
44526_at			AA191741	3.4	3	5.8
51175_at			W52756	3.7	3	5.7
44657_at			AI970788	9	3	0
71526_f_at			AI686949	3	3	1.4
65659_at			AA725644	3.5	3	2.7
44603_at			AA029506	0	3	0
65795_r_at			AL042362	9.4	3	4.7
53416_at			AI341332	7.4	3	6.2
52575_at			H66727	3.3	3	3.6
45208_at			R46263	0.8	3	1.1
74323_at	SUMO-1 activating enzyme subunit 2		AA974493	3.4	3	3.4
52090_at			AI198126	3.9	3	0
62118_at			AI806905	0	3	0.6
51923_at	sphingosine kinase 1	SPHK1	AI769914	14.8	3	1.6
55655_s_at			AI830608	2.6	3	1.9
57226_at	hypothetical protein FLJ22174		AI932981	8	3	0
57280_f_at	major histocompatibility complex, class I, B	HLA-B	AI985880	3.6	3	2.1
58000_at			AA421622	1.2	3	0.9
75213_at			H86747	0	3	2

Figure 15GGGG

67220_at				AW005511	5.3	3	3
41749_at	ES1 (zebrafish) protein, human homolog of	C21ORF33		U53003	4.6	3	4.1
41278_at	BAF53			AF041474	3.8	3	3.3
36792_at	tropomyosin 1 (alpha)	TPM1		Z24727	3	3	2.9
479_at	disabled (Drosophila) homolog 2 (mitogen-responsive phosphoprotein)	DAB2		U53446	3.3	3	3.4
38417_at	adenosine monophosphate deaminase 2 (isoform L)	AMPD2		M91029	3.3	3	2.9
57346_at	Mad4 homolog			AA831438	2.5	3	3.6
52896_at	translocase of inner mitochondrial membrane 10 (yeast) homolog	TIMM10		AI498204	2.2	3	3.7
45276_g_at	protein x 0001			W30943	3.4	3	4.9
45758_at				AA034095	4.6	3	5.9
65612_at	hypothetical protein MGC3180			AW006145	3	3	3
45331_at	hypothetical protein FLJ10355			AA102468	3.2	3	2.7
58647_at	glutathione S-transferase M1	GSTM1		AA203289	4.5	3	3.2
46220_at	peroxisomal short-chain alcohol dehydrogenase			AI131178	3.3	3	3.7
62552_at				AA287801	4.2	3	3.3
49549_at	CD36 antigen (collagen type I receptor, thrombospondin receptor)	CD36		R20784	3	3	2.2
35193_at	chromosome condensation 1-like	CHC1L		AF060219	3.4	2.9	2.2
41295_at	GTT1 protein			AL041780	5.9	2.9	7.4
35247_at	small nuclear RNA activating complex, polypeptide 5, 19kD			AI557062	5.5	2.9	5
41540_at	protein phosphatase 1, regulatory subunit 7	PPP1R7		Z50749	5.2	2.9	5.2
41600_at	proliferation-associated 2G4, 38kD	PA2G4		U59435	3.1	2.9	3
36592_at	prohibitin	PHB		S85655	3.1	2.9	2.1
38424_at	KIAA0747 protein			AB018290	4.3	2.9	4
38770_at	a disintegrin and metalloproteinase domain 12 (meltrin alpha)	ADAM12		AF023476	2.9	2.9	2.7
78442_at	hypothetical protein MGC4692			AI819340	2.9	2.9	2.9
61668_at				AI963090	5	2.9	4.3
54865_at	coronin, actin-binding protein, 1C	CORO1C		N27805	3.2	2.9	3.2
60471_at	hypothetical protein FLJ22439			AA625133	3.2	2.9	3.1
63600_f_at				AW021173	3.1	2.9	3.2
49343_at				AI819630	2.9	2.9	3.2
32800_at	retinoid X receptor, alpha	RXRA		U66306	3	2.9	2.6
91711_at	CDC14 (cell division cycle 14, S. cerevisiae) homolog B	CDC14B		AA173988	3.5	2.9	3.2
72563_at				AI123967	2.3	2.9	2.7

Figure 15HHHH

52932_at	kangai 1 (suppression of tumorigenicity 6, prostate, CD82 antigen (R2 leukocyte antigen, antigen detected by monoclonal and antibody IA4))	KAI1	W51778	3.3	2.9	2.6
45284_at	butyrate-induced transcript 1		AF169958	4.1	2.9	3.8
51163_at	hypothetical protein FLJ22405		AA115361	3.6	2.9	4.5
44558_at	potassium voltage-gated channel, Isk-related family, member 3	KCNE3	AI148745	3.3	2.9	2.9
44849_at	beta-site APP-cleaving enzyme	BACE	AI356651	2.9	2.9	3.7
45805_g_at	hypothetical protein FLJ20643		AA948226	3.1	2.9	3
55040_at	complement component C1q receptor		AA196201	2.8	2.9	2.8
35329_at	cytochrome b5 reductase 1 (B5R.1)		AF091084	3	2.8	2.7
37978_at	quinolinate phosphoribosyltransferase (nicotinate-nucleotide pyrophosphorylase (carboxylating))	QPRT	D78177	2.9	2.8	
38828_s_at	KH-type splicing regulatory protein (FUSE binding protein 2)	KHSRP	AA628946	5.8	2.8	5.8
65813_at	zinc finger protein homologous to Zip-36 in mouse	ZFP36	N32721	3.9	2.8	4.8
54810_i_at	hypothetical protein FLJ10210		AI745170	3	2.8	2.2
48595_at	hypothetical protein FLJ10210		AA707400	3.2	2.8	2.8
72643_i_at	hypothetical protein from EUROMAGE 2021883		AI758570	3.9	2.8	3.2
60863_s_at			AA526910	3.6	2.8	4.3
65465_at			AI953614	2.8	2.8	3.5
59452_at			AI719167	2.8	2.8	2.9
43854_at	xeroderma pigmentosum, complementation group C	XPC	AW004730	3	2.8	2.9
54765_at			AL042673	3.8	2.8	4.2
46569_at			AA552976	3.8	2.8	3.8
54158_at			AI453531	4.3	2.8	4.6
43991_at	hypothetical protein FLJ13576		W19970	3.9	2.8	3.9
38763_at	sorbitol dehydrogenase	SORD	L29254	1.6	2.8	3.1
36642_at	ribosomal protein L27	RPL27	J00287	3.4	2.8	3.9
39830_at	myelin-associated oligodendrocyte basic protein	MOBP	AA044823	3	2.7	5.8
38499_s_at	neurogranin (protein kinase C substrate, RC3)	NRGN	D28113	2.7	2.7	2.7
33925_at	KIAA0982 protein		X99076	2.7	2.7	2.7
35199_at	KIAA0602 protein	XPNPEP2	AB023199	4.3	2.7	4.4
34406_at	X-prolyl aminopeptidase (aminopeptidase P) 2, membrane-bound		AB011174	4.9	2.7	4.6
40296_at	nucleolar protein p40; homolog of yeast EBNA1-binding protein		AL023653	3.2	2.7	3.2
36135_at	topoisomerase (DNA) II beta (180kD)	TOP2B	U86602	2.9	2.7	2.9
36571_at			X68060	3.6	2.7	5.1

Figure 15III

40695_at	IMP (inosine monophosphate) dehydrogenase 1	IMPDH1	J05272	2.9	2.7	3.7
40745_at	adaptor-related protein complex 1, beta 1 subunit	APIB1	L13939	3.4	2.7	3.2
38812_at	laminin, beta 2 (laminin S)	LAMB2	X79683	3	2.7	2.7
56968_at	MADS box transcription enhancer factor 2, polypeptide D (myocyte enhancer factor 2D)	MEF2D	AL118605	4.2	2.7	3.7
81193_r_at			AA884461	2.2	2.7	0.9
48445_at			AA034085	2	2.7	2.6
43926_at			N36085	3.2	2.7	3.2
45335_at	CGI-28 protein	AI037884	AI037884	4.1	2.7	4.4
46890_at			AA203361	2.7	2.7	2.3
54093_at	hypothetical protein		AI307252	3.2	2.7	2.4
61119_at	CCAAT/enhancer binding protein (C/EBP), alpha	CEBPA	AI971171	3.2	2.7	3.1
55671_at	FH1/FH2 domain-containing protein		N90883	2.9	2.7	2.8
40244_s_at	hypothetical protein FLJ11585		AI743654	3.4	2.6	3.5
41209_at	lipoprotein lipase	LPL	M15856	2.8	2.6	2.3
40955_at	prostate tumor over expressed gene 1	PTOV1	U79287	5.2	2.6	4.6
1676_s_at	eukaryotic translation elongation factor 1 gamma	EEF1G	M55409	2.8	2.6	4.6
32455_s_at	proline and glutamic acid rich nuclear protein		U88153	3.3	2.6	3.2
35261_at	glia maturation factor, gamma	GMFG	W07033	3.9	2.6	4.3
36198_at	translocase of outer mitochondrial membrane 20 (yeast) homolog		D13641	6.1	2.6	5.7
38290_at	regulator of G-protein signalling 14	RGS14	AF037195	2.9	2.6	2.7
56426_at	hypothetical protein FLJ22351		AA177127	3	2.6	2.6
59504_at			AA932206	2.6	2.6	2.5
49349_at	PP2135 protein		W72919	2.6	2.6	2.7
90257_at	hypothetical protein DKFZp547D065		AI089323	3.6	2.6	3.7
58235_at	hypothetical protein FLJ10540		AI674163	2.6	2.6	2.5
44638_at	insulin-degrading enzyme	IDE	AA573292	5.2	2.6	4.2
84903_f_at			AI264299	2.8	2.6	2.6
85006_at			AA601045	2.6	2.6	2.6
59095_at			N36764	2.6	2.6	2.4
87624_at			AA668464	0.1	2.6	0.6
864_at	homeo box HB9	HLXB9	U07664	3	2.6	1.9
57918_at	DKFZP586G1517 protein		AA779617	2.8	2.6	2.8
64410_at	KIAA1150 protein		AA813014	4	2.6	3.6
47101_g_at	hypothetical protein		AI972460	3.4	2.6	3.4

Figure 15JJJ

53956_at				AA449296	3.2	2.6	2.8
64890_at	hypothetical protein FLJ10669			AL041174	2.9	2.6	3
46262_at				AA532392	2.7	2.6	3.3
57222_at				AA203532	3.9	2.6	3.5
43499_at	tropomyosin 1 (alpha)	TPM1		N76371	3	2.6	2.6
1933_g_at	ATP-binding cassette, sub-family C (CFTR/MRP), member 5	ABCC5		U83661	3.1	2.5	3
37497_at	hematopoietically expressed homeobox	HHEX		L16499	2.6	2.5	1.6
35519_at				AL049431	3.6	2.5	3.5
34936_at	solute carrier family 4, sodium bicarbonate cotransporter, member 7	SLC4A7		AB012130	2.6	2.5	3.2
39729_at	peroxiredoxin 2	PRDX2		L19185	3.4	2.5	3.3
40824_at	RAN binding protein 16	RANBP16		AB018288	4.8	2.5	5.3
39248_at	aquaporin 3	AQP3		N74607	2.5	2.5	2.4
36322_at	fucosyltransferase 7 (alpha (1,3) fucosyltransferase)	FUT7		AB012668	2.8	2.5	1.4
35342_at				AF052159	2.9	2.5	2.6
32594_at	chaperonin containing TCP1, subunit 4 (delta)	CCT4		AF026291	5.1	2.5	1.7
38336_at	KIAA1013 protein			AB023230	3	2.5	2.9
49902_at				AI927646	1.9	2.5	2.5
58498_at	KIAA1378 protein			AI082236	3.4	2.5	3.4
53721_at				AI742931	2.6	2.5	2.7
88767_at				AW008519	2.5	2.5	2.5
90610_at	leucine-rich neuronal protein	LRN		AI654857	3.3	2.5	4.3
80889_at				AA604308	3.7	2.5	3.3
63886_at	hypothetical protein MGC2616			AI077662	2.9	2.5	3
62273_at				AI766808	2.5	2.5	2.3
91825_at				AI375222	2.5	2.5	2.5
60493_at				AI344328	2.7	2.5	2.6
50198_at				AI208272	2.5	2.5	2.5
44065_at				AI937468	2.5	2.5	2.4
63259_s_at	docking protein 2, 56kD	DOK2		AA864400	2.5	2.5	2.4
61289_at	ATPase, Ca++ transporting, plasma membrane 1	ATP2B1		AI337321	2.5	2.5	2.8
47627_at	cytochrome P540, subfamily IIS, polypeptide 1			AI445492	2.5	2.5	2.5

Figure 15KKKK

affyID	Gene Name	Gene Symbol	GBA	Score_Ecoli	Score_Candida	Score_Influenza
74566_at	insulin-like growth factor-binding protein 4	IGFBP4	A1440424	13.7	75.4	14.1
41045_at	secreted and transmembrane 1	SECTM1	U77643	37.2	67.4	10.4
40314_at	synaptogyrin 3	SYNGR3	AJ002309	2.7	39.7	0.9
39710_at	P311 protein		U30521	6.6	35	8.7
37220_at	Fc fragment of IgG, high affinity Ia, receptor for (CD64)	FCGR1A	M63835	0.5	33.7	2.3
39271_at		RAB4B	AA461365	7.6	33.2	1.5
79210_at			W58117	2.6	28.1	0.2
62804_at			AA621888	0.4	22.9	0.2
75510_at			A1821517	0	21.7	3.2
81915_at	cytotoxic T-lymphocyte-associated protein 4	CTLA4	A1733018	0	19.5	9.8
53670_at			AA664156	4.7	17.4	9.1
32200_at	acid phosphatase, prostate	ACPP	M24902	1.5	15.8	0.4
62743_at	Wiskott-Aldrich syndrome protein interacting protein	WASPIP	AA811088	1.9	15.6	0
34474_at			W28429	1.1	14.2	0.4
64397_at	UCC1 protein		AA131797	2.1	12.7	1.6
41123_s_at	ectonucleotide pyrophosphatase/phosphodiesterase 2 (autotaxin)	ENPP2	L35594	1	12.6	1.8
79822_g_at			A1223310	1.6	11.7	2.4
64344_at	lysosomal-associated membrane protein 1	LAMP1	A1984264	0.8	11.6	0.3
77887_at			AL037345	0	10.8	0
33124_at	vaccinia related kinase 2	VRK2	AB000450	1.3	10.4	2.7
60839_at			A1924996	0.2	10	0
1746_s_at	TNF receptor-associated factor 2	TRAF2	U12597	6.3	9.7	0
36149_at	dihydropyrimidinase-like 3	DPYSL3	D78014	2.3	9.6	1.2
57697_at			A1445628	0	8.8	0.8
1617_at	zinc finger protein 147 (estrogen-responsive finger protein)	ZNF147	D21205	1.8	8.7	0.7
65774_at			A1556963	0.4	8.6	0
58288_at			W63676	2.3	8.5	1.9
56458_at			Z99408	0	8.1	0.8
90732_at	splicing factor, arginine/serine-rich 5	SFRS5	W45137	2.1	8.1	3
36493_at	lymphocyte-specific protein 1	LSP1	M33552	0	8	0
78264_at			N49445	1.7	7.9	1.4
47055_at	hypothetical protein FLJ12287 similar to semaphorins		AA481246	1.1	7.8	0

Figure 16A

33977_at	growth factor independent 1	GFI1	U67369	0.9	7.4	0.9
85241_at			AA846371	1	7.4	0
33524_at	v-crk avian sarcoma virus CT10 oncogene homolog-like	CRKL	X59656	1	7.3	0.4
35704_at	similar to rat HREV107		X92814	2.5	7	0.5
33253_at	KIAA0129 gene product		D50919	0	6.9	0.1
58774_at	eukaryotic translation initiation factor 4 gamma, 1	EIF4G1	AI380979	0.7	6.8	0
82402_at	hypothetical protein DKFZp761D1823		AA522440	0	6.6	0
46246_at	N-Acetylglucosamine kinase		AW026312	0	6.6	0.7
980_at	Niemann-Pick disease, type C1	NPC1	AF002020	0.4	6.5	0
56428_at	hematological and neurological expressed 1	HN1	AI525822	1	6.5	0.7
31859_at	matrix metalloproteinase 9 (gelatinase B, 92kD gelatinase, 92kD type IV collagenase)	MMP9	J05070	1	6.1	0.4
64304_r_at			AW025092	0.6	6	0.8
37068_at	phospholipase A2, group VII (platelet-activating factor acetylhydrolase, plasma)	PLA2G7	U24577	0	5.8	0
76600_at			AI631895	0.5	5.8	0.2
53107_at			N20928	0.1	5.8	0
46497_at			AI524095	0	5.7	0.4
73238_at	collagen, type IX, alpha 1	COL9A1	AI457905	0.6	5.7	0.8
77572_at	hypothetical protein FLJ20429		AI740961	0.1	5.7	0.4
64036_at	hypothetical protein FLJ10716		AI968206	0	5.7	0
39424_at	tumor necrosis factor receptor superfamily, member 14 (herpesvirus entry mediator)	TNFRSF14	U70321	0.2	5.6	0
65082_at	ficollin (collagen/fibrinogen domain-containing) 3 (Hakata antigen)	FCN3	AI150905	0	5.6	0.6
83558_i_at			AI809148	0.9	5.6	0
57617_at			AW009562	1	5.5	1
46601_at	ubiquinol-cytochrome c reductase hinge protein	UQCRH	T16388	1.8	5.5	0
80259_at	hypothetical protein FLJ13171		AA079075	0	5.4	0.4
1526_i_at	fibroblast growth factor 8 (androgen-induced)	FGF8	U47011	0	5.4	0.4
65130_at			AI749588	0.1	5.3	0
69007_r_at			AI583317	1.8	5.3	0.8
41430_at	WD repeat domain 7	WDR7	AB011113	0	5.2	0
39982_r_at	macrophage scavenger receptor 1	MSR1	D13265	0	5.2	0.6
40364_at	solute carrier family 31 (copper transporters), member 1	SLC31A1	U83460	0	5.1	0.8
45110_at			AI989567	0	5.1	0.2

Figure 16B

53718_at	hepatocellular carcinoma-associated antigen 112		AI684645	0.6	5	0.1
41950_at			AA058314	0.7	5	0
90148_at	putative ATPase		AI694453	0	4.9	0.1
44031_at			AI478812	0.9	4.9	0
59617_at	RNA polymerase I transcription factor RRN3		AA830682	0	4.8	0
36372_at	hexokinase 3 (white cell)	HK3	U51333	0	4.7	0
33342_at	RNA, U transporter 1	RNUT1	AF039029	0.4	4.6	0.5
37914_at	KIAA0305 gene product		AB002303	0	4.6	0
54479_at			AA214559	0	4.6	0
61440_at			AA448535	0	4.6	0
32963_s_at	hypothetical GTP-binding protein DKFZp761H171		W27549	0	4.5	0
32574_at	sphingomyelin phosphodiesterase 1, acid lysosomal (acid sphingomyelinase)	SMPD1	X59960	0	4.5	0.1
64369_s_at			AA219354	0.7	4.5	0
49967_at			AI658816	0.4	4.5	0
48055_at			N66640	0	4.4	0
49684_g_at	splicing factor, arginine/serine-rich 11	SFRS11	AI306386	0.7	4.4	0.9
43961_s_at	hypothetical GTP-binding protein DKFZp761H171		AI627648	0.3	4.4	0
44556_at	KIAA1344 protein		AA113931	0	4.4	0.4
65114_at			AA490685	0.1	4.4	0
81793_r_at			AI591416	0.1	4.4	0.5
89975_r_at	CASP8 associated protein 2	CASP8AP2	AI811300	0.4	4.3	0
57613_at	matrix metalloproteinase 25	MMP25	W72721	0	4.3	0
75259_at	complement component 2	C2	AW014901	0	4.3	1.8
72601_at	G-substrate		AI802033	0.8	4.3	0
78478_r_at	FSHD region gene 1	FRG1	H89231	0.6	4.2	0.1
49430_at	g20 protein		AI928807	0	4.2	0
90190_at			AI819924	0	4.2	0
49085_at	vacuolar protein sorting 29 (yeast homolog)	VPS29	AI346485	0.7	4.2	0
77849_r_at			AI499214	0.4	4.1	0.1
84315_at			AA602942	0.1	4.1	0.8
54441_at			AI686323	1	4.1	0.4
54596_at			AI768162	0.5	4.1	0.3
52912_at			3096753	0	4.1	0.2
61842_at	hypothetical protein FLJ20716		AI769900	1	4	0.8

Figure 16C

75244_at					AI732578	0.8	3.9	0.4
56279_at	GTF2I repeat domain-containing 1		GTF2IRD1		R12398	0.2	3.9	0
52164_at	chromosome 11 open reading frame 24		C11ORF24		AA065185	0	3.9	0
44580_at	pregnancy-induced growth inhibitor				AI814233	0	3.9	0
34128_at	UDP-Gal:betaGlcNAc beta 1,3-galactosyltransferase, polypeptide 5		B3GALT5		AB020337	0	3.8	0.3
39298_at	alpha2,3-sialyltransferase				AB022918	0	3.8	0
34864_at	hypothetical protein				AF070638	0.9	3.8	0
36958_at	zyxin		ZYX		X95735	0	3.8	0
38172_at	carbonyl reductase 1		CBR1		AB003151	0	3.8	0
41057_at	uncharacterized hypothalamus protein HT012				AI760162	0	3.8	0
57037_at	kinesin family member 5B		KIF5B		AA057424	0	3.8	0.9
69879_at					AI688187	0.2	3.8	0
60823_at					AA191336	2.6	3.8	0.2
75262_s_at					AW025934	0.2	3.8	0
48413_at	ras GTPase activating protein-like				W46352	0.6	3.8	0
32079_at	kinesin family member 13B		KIF13B		AB014539	0.3	3.7	0.6
77696_at					AI733058	0	3.7	0.3
35820_at	GM2 ganglioside activator protein		GM2A		X62078	0.6	3.7	0
82660_at	ninein (GSK3B interacting protein)		NIN		T82318	0	3.7	0.2
40038_at	suppression of tumorigenicity 7		ST7		W02490	0.4	3.6	2.4
38373_g_at					U66042	0.7	3.6	0
58872_at					AI983986	0	3.6	0.9
70734_at					R14439	0.7	3.6	0.5
73198_at					AL043551	0	3.6	0.5
52608_at					AI860012	0.3	3.6	0
62976_at					AI251931	0.6	3.6	0
81555_at	hypothetical protein FLJ20733				AI360255	0	3.5	0.4
46958_at					AI868421	0	3.5	0.9
65800_s_at	hypothetical protein				T89723	0.4	3.5	0.8
72999_f_at	chromosome 8 open reading frame 1		C8ORF1		AW002557	0.9	3.5	0.4
38223_at	vascular Rab-GAP/TBC-containing				AB024057	1	3.4	0.5
37392_at	phosphorylase kinase, beta		PHKB		X84908	0.9	3.4	0
57990_g_at					AI206946	0	3.4	0
49935_at					AI886666	0.2	3.4	0.4

Figure 16D

64903_at				N30209	0	3.4	0.4
84263_at				AA525944	0.8	3.4	0
85274_at				AI939617	1	3.4	0.3
67519_at				AI498953	0.5	3.4	0.5
74738_r_at	RNA binding motif, single stranded interacting protein 1		RBMS1	AW007869	0.4	3.4	0.2
81533_at				AI766311	0	3.4	0.2
51902_at	hypothetical protein			AA215376	0	3.4	0
37737_at	protein-L-isoaspartate (D-aspartate) O-methyltransferase		PCMT1	D25547	0	3.3	0
74772_at				AI424825	0.4	3.3	0.5
78695_at				AW001771	0	3.3	0
89164_at				AI393930	1	3.3	0.5
90033_at	phosphodiesterase 2A, cGMP-stimulated		PDE2A	T66157	0.6	3.3	0.1
47508_at	centaurin-alpha 2 protein			AW003904	0.5	3.3	0.5
45919_at				AA021104	0.5	3.3	0.4
88448_at	hypothetical protein FLJ11113			AI911379	0	3.3	0
52641_at	putative G protein-coupled receptor			AI828579	0.1	3.3	0.4
32477_at	calpain 9 (nCL-4)		CAPN9	AF022799	0.2	3.2	0.7
1499_at	farnesyltransferase, CAAX box, alpha		FNTA	L10413	0	3.2	0.7
41225_at	dual specificity phosphatase 3 (vaccinia virus phosphatase VH1-related)		DUSP3	AL049417	0.1	3.2	0
34196_at				AI337901	0.2	3.2	0
32544_s_at	Ras suppressor protein 1		RSU1	L12535	0.6	3.2	0
76912_at				AI081271	0	3.2	0.1
65440_at	SCO (cytochrome oxidase deficient, yeast) homolog 1		SCO1	AA205413	0.9	3.2	0
47911_at				AI249173	0	3.2	0
68708_at				AI560174	0.8	3.2	0.5
60791_at	hypothetical protein FLJ11011			AI828160	0	3.2	0
62487_i_at				W51746	0.3	3.2	0
61277_at	peptide transporter 3			AI141684	0.7	3.2	0
31742_at	tumor necrosis factor (ligand) superfamily, member 14		TNFSF14	AF064090	0.6	3.1	0.1
39778_at	mannosyl (alpha-1,3-)-glycoprotein beta 1,2-N-acetylglucosaminyltransferase		MGAT1	M55621	0.4	3.1	0.5
949_s_at	proteasome (prosome, macropain) 26S subunit, ATPase, 6		PSMC6	D78275	0.5	3.1	0.2
87791_at				AA670223	0	3.1	0
74870_at	carnitine palmitoyltransferase I, liver		CPT1A	AI675101	0.3	3.1	0.4

Figure 16E

54622_at	receptor-interacting serine-threonine kinase 3	RIPK3	N99896	0.9	3.1	0
43952_at			A1721219	0.3	3.1	0
61712_at	hypothetical protein PRO0823		A1990512	0.4	3.1	0
51857_at			AA705016	0.6	3.1	0
57217_at	biphenyl hydrolase-like (serine hydrolase; breast epithelial mucin-associated antigen)	BPHL	AA526961	0	3.1	0
41710_at	hypothetical protein		AL079277	0.7	3	0
1399_at	transcription elongation factor B (SIII), polypeptide 1 (15kD, elongin C)	TCEB1	L34587	0.3	3	0
38944_at	MAD (mothers against decapentaplegic, Drosophila) homolog 3	MADH3	U68019	0	3	0.4
90346_at			AW003138	0	3	0
44632_at			AA007697	0	3	0.1
75317_s_at	interferon gamma receptor 2 (interferon gamma transducer 1)	IFNGR2	A1985833	0.9	3	0
53888_at	cathepsin B	CTSB	R81090	0	3	0
44603_at			AA029506	0	3	0
62118_at			A1806905	0	3	0.6

Figure 16F

affyID	Gene Name	Gene Symbol	GBA	Score_Ecoli	Score_Candida	Score_Influenza
1520_s_at	endothelin 1	EDN1	J05008	4444	283.5	113.7
1369_s_at	interleukin 8	IL8	M28130	2300.5	32.6	122.1
35822_at	B-factor, properdin	BF	L15702	1635.7	287.1	17.3
38299_at	interleukin 6 (interferon, beta 2)	IL6	X04430	1566.4	101.3	227.7
40385_at	small inducible cytokine subfamily A (Cys-Cys), member 20	SCYA20	U64197	943.4	28.3	53.8
91636_at			A1652725	886.7	27.7	7
1491_at	pentaxin-related gene, rapidly induced by IL-1 beta	PTX3	M31166	827.3	8.9	79
37187_at	GRO2 oncogene	GRO2	M36820	785	32	26.1
35061_at	small inducible cytokine subfamily B (Cys-X-Cys), member 11	SCYB11	AF030514	655.1	844.8	729.7
1372_at	tumor necrosis factor, alpha-induced protein 6	TNFAIP6	M31165	573.9	37.7	75.1
425_at	interferon, alpha-inducible protein 27	IFI27	X67325	565.9	754	421
2002_s_at	BCL2-related protein A1	BCL2A1	U27467	547.8	65.2	70.5
37944_at	GTP cyclohydrolase 1 (dopa-responsive dystonia)	GCH1	U19523	505.4	330.9	169
1069_at	prostaglandin-endoperoxide synthase 2 (prostaglandin G/H synthase and cyclooxygenase)	PTGS2	U04636	495.3	26.7	53.1
77706_at			AA641972	493	59	7.8
408_at	GRO1 oncogene (melanoma growth stimulating activity, alpha)	GRO1	X54489	486.7	11.5	7.1
38549_at			AF026941	484.1	612.9	1149.5
39402_at	interleukin 1, beta	IL1B	M15330	468.5	30.5	12.2
52615_at			AA948319	425	188.4	94.9
33304_at	interferon stimulated gene (20kD)	ISG20	U88964	327.2	200	103.7
909_g_at	interferon-induced protein with tetratricopeptide repeats 2	IFIT2	M14660	320.5	145.5	573.3
43853_at	hypothetical protein		AA618602	296.2	15.3	18.6
1021_at	interferon, gamma	IFNG	J00219	289.9	58.8	36.2
41677_at	interleukin 15 receptor, alpha	IL15RA	AF035279	278	99.5	111.4
1097_s_at	chemokine (C-C motif) receptor 7	CCR7	L31584	276.6	100.6	88.3
37219_at	monokine induced by gamma interferon	MIG	X72755	261.6	704.2	182.7
33955_at	interleukin 12A (natural killer cell stimulatory factor 1, cytotoxic lymphocyte maturation factor 1, p35)	IL12A	M65291	250.2	18.4	14.6
39827_at	hypothetical protein		AA522530	245.1	38.4	14.6

Figure 17A

33705_at	phosphodiesterase 4B, cAMP-specific (dunce (Drosophila)-homolog phosphodiesterase E4)	PDE4B	L20971	232.3	51.8	92.7
1403_s_at	small inducible cytokine A5 (RANTES)	SCYA5	M21121	232.1	90.6	200.3
48864_at	interferon, alpha-inducible protein 27	IFI27	AI991845	216.4	274.9	141.4
36116_at	thyroid hormone receptor interactor 10	TRIP10	AJ000414	215.9	56.7	7
61371_at	cyclin-E binding protein 1		AA555023	204	148	479.5
1852_at	tumor necrosis factor (TNF superfamily, member 2)	TNF	X02910	201.2	47.5	57
45164_at	19A24 protein		W74027	189.7	74.5	32.5
62330_at			AI075407	185.9	169.1	121.9
49148_at			AA947472	182.2	52	423.6
38584_at	interferon-induced protein with tetratricopeptide repeats 4	IFIT4	AF026939	179.4	153.2	148.6
58918_at			AA210892	176.1	24	34.8
39959_at	gamma-aminobutyric acid (GABA) B receptor, 1	GABBR1	AL031983	161.4	263.9	33.7
57761_at			AI313253	161	50.6	33.7
40159_r_at	neutrophil cytosolic factor 1 (47kD, chronic granulomatous disease, autosomal 1)	NCF1	M55067	160.7	91	140.3
31623_f_at	metallothionein 1A (functional)	MT1A	K01383	160.3	22.3	49.7
40323_at	CD38 antigen (p45)	CD38	D84276	160.1	102.8	18.6
41048_at	phorbol-12-myristate-13-acetate-induced protein 1	PMAIP1	D90070	159.8	9.8	408.5
1107_s_at	interferon-stimulated protein, 15 kDa		M13755	159	86.3	213.8
36296_at	lymphotoxin alpha (TNF superfamily, member 1)	LTA	D12614	154	62.8	28.5
431_at	small inducible cytokine subfamily B (Cys-X-Cys), member 10	SCYB10	X02530	152	390	545.7
59215_at	RNA helicase		AI807018	151.1	63	36.2
34666_at	superoxide dismutase 2, mitochondrial	SOD2	X07834	141.3	63.1	42.3
89884_at	metallothionein 2A	MT2A	AI991852	138	45.1	82.3
32114_s_at	adenosine A2a receptor	ADORA2A	S46950	137.9	31	27.9
60896_at	integrin, beta 8	ITGB8	AI634852	137.5	31.7	10.7
38754_at	p8 protein (candidate of metastasis 1)		AI557295	131.1	35.6	632.5
48919_at	potassium inwardly-rectifying channel, subfamily J, member 2	KCNJ2	AA393850	130	11.2	13.1
51725_at			AA521246	129.3	35.4	18
63335_at	sorting nexin 10		AI285531	127.4	55.8	74.2
48753_at	PPAR(gamma) angiopoietin related protein		N57259	126	7.9	14.7
53508_at			AI247103	124.7	84.6	27.2

Figure 17B

63628_at	interleukin 7 receptor	IL7R	AI655781	123.1	41.5	20.9
1548_s_at	interleukin 10	IL10	U16720	121.9	4.3	3.2
70458_i_at			AI654525	120.3	174.3	127.2
36174_at	macrophage myristoylated alanine-rich C kinase substrate		X70326	119.6	44.8	42
84995_at			AA769482	117.4	12.2	6.6
50978_at	putative secreted ligand homologous to ffx1	hfx	AI338625	117.2	3.8	3.3
609_f_at	metallothionein 1B (functional)	MT1B	M13485	116.3	11.1	20.1
51972_at	ubiquitin specific protease 18	USP18	AA143794	114.4	158.3	218
51546_at			AI498361	113.1	77.9	73.9
63066_at			AA001735	113	12.6	9.6
675_at	interferon induced transmembrane protein 1 (9-27)	IFITM1	J04164	112.7	140.9	173.6
71839_at			AA251131	111.4	76.5	82.8
58957_at	hypothetical protein FLJ20637		AI620475	108.9	42.7	28.3
65867_at	FXYD domain-containing ion transport regulator 6	FXYD6	AL043089	103.8	73.3	28.2
1370_at	interleukin 7 receptor	IL7R	M29696	102.3	30.8	25.3
31622_f_at	metallothionein 1F (functional)	MT1F	M10943	101.8	13.1	32.6
70097_at	indoleamine-pyrrole 2,3 dioxygenase	INDO	AI302268	99.5	29.8	39.9
59777_at			AA766831	98.3	63.3	41.1
65626_at			AA059458	96.4	20.4	9
48035_at	A kinase (PRKA) anchor protein 2	AKAP2	AA101125	95.8	38.5	43.7
88714_at			AA873008	94.5	63.3	12.9
76710_at	cardiotrophin-like cytokine; neurotrophin-1/B-cell stimulating factor-3		AI040033	92.7	4.7	4
39081_at	metallothionein 2A	MT2A	AI547258	92.5	37.3	94.1
44815_s_at			AA733119	92	11.1	28.6
39594_f_at	metallothionein 1H	MT1H	R93527	91.6	9.4	26.1
36804_at	indoleamine-pyrrole 2,3 dioxygenase	INDO	M34455	91.3	36.9	37.7
56246_at	6-phosphofructo-2-kinase/fructose-2,6-biphosphatase 3	PFKFB3	AI765775	91	6.2	11.3
37168_at	similar to lysosome-associated membrane glycoprotein		AB013924	90.9	50.6	16.1
870_f_at	metallothionein 3 (growth inhibitory factor (neurotrophic))	MT3	M93311	90.7	15.1	42.9
46659_at			AI742057	87.7	54.8	56
57085_s_at	DKFZP586G0522 protein		AW008790	85.4	4.8	52.3
54982_at	pellino (Drosophila) homolog 1	PELL1	AL043980	85.4	6.2	54.5
45237_at	superoxide dismutase 2, mitochondrial	SOD2	AA142976	84.4	32.6	17.7

Figure 17C

38570_at	major histocompatibility complex, class II, DO beta	HLA-DOB	X03066	84.2	55.5	30.1
91313_at			AA131041	84	77.5	157.1
84893_at			AI446168	83.4	119.5	160.2
1527_s_at			U50527	82.5	50.7	14.2
32775_r_at	phospholipid scramblase 1	PLSCR1	AB006746	81.8	68.4	32.6
36130_f_at	metallothionein 1E (functional)	MT1E	R92331	78.7	4.6	8.6
54594_at	KIAA0668 protein		AA009571	78.6	23.3	31.3
35844_at	syndecan 4 (amphiglycan, ryudocan)	SDC4	D79206	77.4	5.8	20.7
44126_at			W28501	73.5	28.6	16.2
69142_at			AI635522	72.4	7.8	8.6
31540_at	tumor necrosis factor receptor superfamily, member 9	TNFRSF9	U03397	72.2	7.8	19.5
35735_at	guanylate binding protein 1, interferon-inducible, 67kD	GBP1	M55542	72.1	49.4	49.8
39224_at	KIAA0580 protein		AB011152	71.4	27.4	18
41446_f_at	RNA helicase-related protein		H68340	71.4	10.1	21.9
38111_at	chondroitin sulfate proteoglycan 2 (versican)	CSPG2	X15998	71.2	4.1	27.4
47483_at	similar to rat myomegalin		AI659612	71.1	9.8	11.8
50746_at			AA897516	71	20.1	43.9
73455_f_at			H72723	70.4	8.2	8.5
37643_at	tumor necrosis factor receptor superfamily, member 6	TNFRSF6	X63717	69	17.4	30
44513_at			AI760613	68.8	81.5	36.6
39114_at	decidual protein induced by progesterone		AB022718	68.6	26.6	28.6
64444_at	hypothetical protein		AI800563	68.6	84.1	136.7
37641_at	interferon-induced, hepatitis C-associated microtubular aggregate protein (44kD)		D28915	68.5	51.1	43.6
49438_at						
32026_s_at	PDZ domain containing guanine nucleotide exchange factor(GEF)1		AI535730	68.4	21	15.8
			AB002311	68.3	25.1	16.6
63769_at			AI749656	67.8	49	14.3
71174_r_at	KIAA1682 protein		AI921158	67.8	81.9	115.6
1715_at	tumor necrosis factor (ligand) superfamily, member 10	TNFSF10	U37518	66.8	179.7	463.3
41475_at	ninjurin 1	NINJ1	U91512	65.2	35.7	16.7
595_at	tumor necrosis factor, alpha-induced protein 3	TNFAIP3	M59465	65.1	11.4	26.6
1036_at	interleukin 15	IL15	U14407	64.7	16.6	22.1
77244_at	bHLH factor Hes4		AI380603	64.7	15.4	23.6
286_at	H2A histone family, member O	H2AFO	L19779	64.6	21	57.1

Figure 17D

38326_at	putative lymphocyte G0/G1 switch gene		M69199	64.6	20.8	6.5
51015_at			AI655467	64.4	21.6	4.4
75014_i_at	KIAA1533 protein		AW006583	64.4	37.7	3.6
73236_g_at	guanylate binding protein 1, interferon-inducible, 67kD	GBP1	AW014593	63.9	55.8	29.8
38631_at	tumor necrosis factor, alpha-induced protein 2	TNFAIP2	M92357	63.8	17.8	18.4
36227_at			AFO43129	63.4	21.8	18.5
1717_s_at	baculoviral IAP repeat-containing 3	BIRC3	U45878	63.1	47.7	20.2
78482_at			AA705165	62.7	3.6	35.7
39221_at	leukocyte immunoglobulin-like receptor, subfamily B (with TM and ITIM domains), member 2	LILRB2	AFO04231	62.6	6.1	17.8
87102_at			AI823649	62	51.5	48.7
51920_at	melanoma differentiation associated protein-5		AA134958	61.8	41.7	22.4
45289_at	novel Ras family protein		AA133248	61.7	3.8	24.3
49140_at			AI244908	61.5	3.8	4.4
879_at	myxovirus (influenza) resistance 2, homolog of murine CD80 antigen (CD28 antigen ligand 1, B7-1 antigen)	MX2	M30818	61.1	61	69
35015_at		CD80	M27533	61	26.5	15.7
44981_s_at			AA115266	59.2	45	5.1
38121_at	tryptophanyl-tRNA synthetase	WARS	X59892	58.2	45.4	45.6
1914_at	cyclin A1	CCNA1	U66838	57.6	16	70.3
67713_r_at			AI986192	57.5	31.9	4.2
32606_at	brain acid-soluble protein 1	BASP1	AA135683	56.9	17.8	11
37163_at	DKFZP586C1619 protein		AL050374	56.3	9.1	10.6
32700_at	guanylate binding protein 2, interferon-inducible	GBP2	M55543	55.5	23.9	6.1
1867_at	CASP8 and FADD-like apoptosis regulator	CFLAR	AF005775	54.9	24.7	5.8
906_at	signal transducer and activator of transcription 4	STAT4	L78440	54.6	10.3	5.9
1377_at	nuclear factor of kappa light polypeptide gene enhancer in B-cells 1 (p105)	NFKB1	M58603	54.5	18.3	18
89899_at	myxovirus (influenza) resistance 2, homolog of murine	MX2	AW001846	54	64.3	70.9
1891_at	mitogen-activated protein kinase kinase 8	MAP3K8	D14497	53.8	4.5	11.2
76762_at			AI246590	53.8	8.9	10.2
37014_at	myxovirus (influenza) resistance 1, homolog of murine (interferon-inducible protein p78)	MX1	M33882	53.6	82.5	69.2
649_s_at	chemokine (C-X-C motif), receptor 4 (fusin)	CXCR4	L06797	53.5	40.7	36
74633_at	tumor necrosis factor, alpha-induced protein 2	TNFAIP2	AI986430	53.5	15.5	6.1
63382_at	interferon consensus sequence binding protein 1	ICSBP1	AI073984	53.5	35.7	8

Figure 17E

36927_at	hypothetical protein, expressed in osteoblast		AB000115	53.4	60.4	83.5
37137_at	granzyme B (granzyme 2, cytotoxic T-lymphocyte-associated serine esterase 1)	GZMB	M17016	52.3	9.1	15.3
80773_at			H15073	51.8	9.8	13.9
48025_at	NCK adaptor protein 2	NCK2	A1934552	51.7	5.9	6.9
63780_at	hypothetical protein FLJ11259		AA814195	51.4	19.6	26.4
62528_at	hypothetical protein FLJ20073		AA741307	51.4	23.4	10.3
222_at	exostoses (multiple) 1	EXT1	S79639	50.9	5	9.8
37451_at			AL109695	50.6	4.8	5.8
35143_at	KIAA1031 protein		AI138605	50.6	24.1	4.6
669_s_at	interferon regulatory factor 1	IRF1	L05072	50.6	35.4	27.2
34435_at	aquaporin 9	AQP9	AB008775	50.5	15.3	3.4
32814_at	interferon-induced protein with tetratricopeptide repeats 1	IFIT1	M24594	50.5	70.1	76.9
41745_at	interferon induced transmembrane protein 3 (1-8U)	IFITM3	X57352	49.6	74	69.9
36412_s_at	interferon regulatory factor 7	IRF7	U53831	49.2	41.5	54.1
35985_at	A kinase (PRKA) anchor protein 2	AKAP2	AB023137	49	20.1	24.6
70460_f_at			AI654525	48.9	90.2	75.7
60103_at			AI916948	46.9	13.9	44.2
41384_at	receptor-interacting serine-threonine kinase 2	RIPK2	AF117829	46.7	26.8	104.9
54565_at			AA149736	45.6	8.7	10.3
51670_i_at			AI161358	45.6	13.2	52.1
529_at	dual specificity phosphatase 5	DUSP5	U15932	45	12.1	10.5
43939_at			AI590710	44.5	9.7	18
53970_at	dual adaptor of phosphotyrosine and 3-phosphoinositides		AF150266	43.8	9.5	3
75246_s_at	solute carrier family 1 (glial high affinity glutamate transporter), member 3	SLC1A3	W26838	43.1	7.2	4.8
40419_at	erythrocyte membrane protein band 7.2 (stomatin)	EPB72	X85116	42.8	19	21.9
56373_at	UDP-Gal:betaGlcNAc beta 1,4- galactosyltransferase, polypeptide 5	B4GALT5	AA133969	42.2	9.3	9.5
43524_at	KIAA1170 protein		C14031	42.1	18.4	7.2
88629_at			AI337136	42	18.5	15.7
42363_r_at	STAT induced STAT inhibitor 3		AI680350	41.7	5.3	5.8
60903_s_at	T-cell lymphoma invasion and metastasis 2	TIAM2	AI126294	41.6	25.5	6.8

Figure 17F

39775_at	serine (or cysteine) proteinase inhibitor, clade G (C1 inhibitor), member 1	SERP1NG1	X54486	41.5	67.8	17.1
75868_r_at						
31832_at	KIAA0286 protein		A1791208	41.5	10.4	7
33849_at	pre-B-cell colony-enhancing factor		AB006624	41.3	5	5.1
34439_at	absent in melanoma 2		U02020	41	8.3	16.3
61671_at		AIM2	AF024714	40.9	18.1	16
44640_at			AA194980	40.7	35.3	10
64333_s_at	pyrophosphatase (inorganic)	PP	W30985	40.2	11.6	8.4
41654_at	adenosine deaminase	ADA	AW009649	39.9	10.9	5
53609_at			X02994	39.7	4.1	3.2
38388_at	2',5'-oligoadenylate synthetase 1 (40-46 kD)	OAS1	C14904	39.3	13	22.8
40357_at	inhibin, beta A (activin A, activin AB alpha polypeptide)	INHBA	M11810	38.7	54	103.7
80915_at	transcription factor ets		J03634	38.6	4.9	5
34745_at			A1347001	38.2	44.7	11.3
51104_at			AF070570	38.1	14.3	8.7
59283_at	hypothetical protein FLJ20035		A1937446	37.7	10.5	22.6
62130_at	28kD interferon responsive protein		AL042790	37.5	24.6	11.4
38576_at	H2B histone family, member B		AA651720	37.5	34	40
50124_at		H2BFB	AJ223353	37.4	13.7	16.1
41045_at	secreted and transmembrane 1	SECTM1	AA580047	37.3	29.6	43.2
47090_at	similar to mouse Xrn1 / Dhm2 protein		U77643	37.2	67.4	10.4
39070_at	singed (Drosophila)-like (sea urchin fascin homolog like)	SNL	A1004667	37.2	45.7	19.5
38389_at	2',5'-oligoadenylate synthetase 1 (40-46 kD)	OAS1	U03057	36.9	23.6	31
90421_at	conserved gene amplified in osteosarcoma		X04371	36.5	71.8	35
69876_at			AA633203	36.5	35.1	8.6
62214_at	hypothetical protein FLJ11354		A1825713	36.1	32.1	60.3
45256_at	pre-B-cell colony-enhancing factor		A1583960	35.5	28.7	23.8
46133_at			A1271460	35.1	7.1	33.4
40153_at	ATP-binding cassette, sub-family B (MDR/TAP), member 2	ABCB2	W60263	34.8	34.9	12.6
37294_at	B-cell translocation gene 1, anti-proliferative	BTG1	X57522	34.3	33	40.6
54769_at			X61123	34.3	15.2	8
55353_at			AA203416	33.9	10.8	14.7
848_at	TNF receptor-associated factor 1	TRAF1	AA760977	33.6	6.9	6
46760_r_at			U19261	33.5	9.9	5.5
			H38110	33.5	9.1	5.8

Figure 17G

52183_at				AA243659	33.5	15.4	18.8
52085_at				AA210833	33.1	21	30.6
36780_at	clusterin (complement lysis inhibitor, SP-40,40, sulfated glycoprotein 2, testosterone-repressed prostate message 2, apolipoprotein J)	CLU		M25915	32.9	5	8.7
59365_at				D59637	32.9	6.9	4.9
56922_at				AA211158	32.7	18.2	35.1
47484_at				H17489	32.5	22.9	8.8
1445_at	chemokine (C-C motif) receptor-like 2	CCRL2		AF014958	32.3	6.4	3.5
1633_g_at	pim-2 oncogene	PIM2		U77735	32	15.7	11.4
41635_at	Wilms' tumour 1-associating protein			D14661	31.6	6.1	18.4
47113_at	Kruppel-like factor 5 (intestinal)	KLF5		AI815057	31.5	41.2	7.2
35254_at	FLN29 gene product			AB007447	31.4	36.2	36.8
81966_at				AI199418	31.2	19.8	24.4
36575_at	regulator of G-protein signalling 1	RGS1		S59049	31	46.9	16.7
33063_at	glycoprotein Ib (platelet), alpha polypeptide	GP1BA		JO2940	30.9	9.1	9.4
36674_at	small inducible cytokine A4 (homologous to mouse Mip-1b)	SCYA4		JO4130	30.3	13.4	13.3
42524_at				R37337	30.1	6.4	3.6
60835_at				AI831561	30.1	9	5.6
58944_s_at	ribosomal protein S6 kinase, 52kD, polypeptide 1	RPS6KC1		AA670286	30	7	5.6
63982_at	ubiquinol-cytochrome c reductase (6.4kD) subunit			AI480328	29.8	8.6	9.8
40505_at	ubiquitin-conjugating enzyme E2L 6	UBE2L6		AA883502	29.7	38.7	20
49339_s_at	reticulocalbin 1, EF-hand calcium binding domain	RCN1		AI817760	29.7	11.9	6.9
84909_at				AA902134	29.7	55.3	39.5
53588_at				AI967984	29.2	4.3	33.3
190_at	nuclear receptor subfamily 4, group A, member 3	NR4A3		U12767	29.1	24.6	25.8
80045_at				AI989871	29.1	26.8	12.8
40362_at	nuclear factor of kappa light polypeptide gene enhancer in B-cells 2 (p49/p100)	NFKB2		X61498	29	11.4	5.2
67764_r_at				AA004879	29	4.4	11.8
44729_at	hypothetical protein FLJ13855			AI027524	28.8	10.5	13.3
53010_at				AI809925	28.6	4.2	29.8
1953_at	vascular endothelial growth factor	VEGF		AF024710	28.4	6.2	7.2
56999_at				AI625959	28.4	11	3.4

Figure 17H

60424_at	hypothetical protein DKFZp761B2423		AA971604	28.4	8.4	58
89202_at			AI347361	28.3	30.4	9.1
58941_at	phosphatidylserine-specific phospholipase A1alpha		AI765967	28.1	24.1	11.5
52337_g_at			AI762208	28	8.5	7.5
81887_at			AI766826	27.8	10.9	33.8
44825_at	RAD50 (S. cerevisiae) homolog	RAD50	AA126482	27.8	17.5	30.9
37442_at			AL050378	27.6	5	11.8
57273_s_at	KIAA0477 gene product		AI634721	27.6	11.9	5.8
883_s_at	pim-1 oncogene	PIM1	M54915	27.5	3.1	4.6
54491_at			AA043562	27.5	3	3.1
38617_at	LIM domain kinase 2	LIMK2	D45906	27.3	4.4	4
34777_at	adrenomedullin	ADM	D14874	27.2	8.5	7.1
44115_at	hypothetical protein from EUROIMAGE 588495		AA486504	27.1	3.1	3.8
87497_f_at			AI832016	27	21.1	7.7
52905_at	Rag C protein		AI279898	26.9	14.5	19.7
39263_at	2'-5' oligoadenylate synthetase 2	OAS2	M87434	26.7	28	34.8
49117_at	interferon-induced protein 75, 52kD	IFI75	AI743445	26.5	24.4	71
38970_s_at	Nef-associated factor 1		AJ011896	26.4	7.6	4.2
40496_at	complement component 1, s subcomponent	C1S	J04080	26.3	5.6	7.9
1737_s_at	insulin-like growth factor-binding protein 4	IGFBP4	M62403	26.2	72	18.6
1173_g_at				26.2	15.6	19.2
64047_at	KIAA1376 protein		AA587245	26.1	6.7	38.4
41744_at	tumor necrosis factor alpha-inducible cellular protein containing leucine zipper domains; Huntingtin interacting protein L; transcription factor IIIA-interacting protein		AF070533	26	45.9	21.1
64733_r_at	serine (or cysteine) proteinase inhibitor, clade B (ovalbumin), member 9	SERPINB9	AA156247	26	12	6.4
48531_at			AA522816	26	9	3.6
75000_at			AI735440	25.9	12.1	6.5
56871_at			AI953847	25.7	8.6	12.8
62992_at			AI214996	25.6	10	4.3
56769_at	B7-H1 protein		AA292201	25.6	8	6.1
60968_at			AI990212	25.3	8.3	4.9
49459_at	apolipoprotein L	APOLI	AA156784	25.2	18.2	17.4
90662_at	2'-5' oligoadenylate synthetase 2	OAS2	AI340262	25.2	27.3	19

Figure 171

64450_at	2'-5'oligoadenylate synthetase 3	OAS3	AA135525	25.2	30.3	10.8
38788_at	promyelocytic leukemia	PML	M82827	25.1	20.3	8.8
33924_at	KIAA1091 protein		AB029014	24.9	10.8	5.1
464_s_at	interferon-induced protein 35	IFI35	U72882	24.3	35.4	12.9
41742_s_at	tumor necrosis factor alpha-inducible cellular protein containing leucine zipper domains; Huntingtin interacting protein L; transcription factor IIIA-interacting protein		AF061034	24.3	28.1	8.7
36175_s_at	human immunodeficiency virus type I enhancer-binding protein 2	HIVEP2	AL023584	24.2	7	5.7
48964_at			AI629027	24.2	17.2	9.1
48076_at	hypothetical protein		AI949434	24.1	7.5	19.9
83874_at			AI024818	23.9	7.8	4.4
46266_at	serine (or cysteine) proteinase inhibitor, clade A (alpha-1 antitrypsin, antitrypsin), member 8	SERPINA8	AI208913	23.9	15.5	9.9
39422_at	KIAA0477 gene product		AB007923	23.8	11.2	7.1
47899_at			AA056755	23.8	16.9	13.4
62217_at			T81422	23.7	7.2	15.8
39768_at	2'-3'-cyclic nucleotide 3' phosphodiesterase	CNP	D13146	23.7	9.9	7.8
53975_at	KIAA1151 protein		Z78399	23.6	3.4	4.7
54744_at			AA582287	23.5	8	3
32156_at	poliovirus receptor-related 2 (herpesvirus entry mediator B)	PVRL2	AF044968	23.4	5.9	8.9
41503_at	KIAA0854 protein		AB020661	23.1	10.5	8.6
41142_at	3-oxoacid CoA transferase	OXCT	U62961	23	3.7	4.5
35718_at	interferon-induced protein 41, 30kD	IFI41	L22342	23	37.3	14.5
41472_at	phorbol-like protein MDS019		AL078641	22.9	41	25.5
47870_r_at	hypothetical protein DKFZp761C169		AW003245	22.9	4	8.8
908_at	interferon-induced protein with tetratricopeptide repeats 2	IFIT2	M14660	22.9	34	384.9
38281_at	caspase 7, apoptosis-related cysteine protease	CASP7	U67319	22.7	14.4	4.9
87415_at			W73083	22.7	4.8	17.3
39119_s_at	natural killer cell transcript 4		AA631972	22.5	34.2	10.5
946_at	t-complex-associated-testis-expressed 1-like 1	TCTEL1	D50663	22.1	20.3	12.3
68002_at			AA503803	21.9	17.2	13.1
32529_at	transmembrane protein (63kD), endoplasmic reticulum/Golgi intermediate compartment		X69910	21.7	23.9	6.8

Figure 17J

58605_at	DNA segment on chromosome X (unique) 9928 expressed sequence	DXS9928E	AI697939	21.5	20.6	9.4
44092_at			AI862887	21.3	17.7	5.5
39660_at	defensin, beta 1	DEFB1	AI309115	21.1	7.8	15.1
51092_at	putative zinc finger protein NY-REN-34 antigen		AI950095	21.1	14.6	8.5
51161_at	ras homolog gene family, member A	ARHA	AI743085	20.9	5.2	20.3
40852_at	tudor repeat associator with PCTAIRE 2		AB025254	20.7	18.9	29.9
64247_at			H40631	20.6	4	3.1
39517_at	hypothetical protein 23851		AF035313	20.5	14.3	16.9
106_at	run-related transcription factor 3	RUNX3	Z35278	20.4	17	11.3
34699_at	CD2-associated protein	CD2AP	AL050105	20.4	12.9	8
40447_at	lipin 2	LPIN2	D87436	20.2	47.2	18.2
91350_at			AA989283	20.2	4.3	3.7
63969_f_at	clusterin (complement lysis inhibitor, SP-40,40, sulfated glycoprotein 2, testosterone-repressed prostate message 2, apolipoprotein J)	CLU	AI982754	20.2	3	3.5
53855_at			N36284	20.1	12.7	12.5
50296_i_at	apolipoprotein L 3	APOL3	T63428	20	24.4	12.1
269_at	2'-5'oligoadenylate synthetase-like	OASL	L40387	19.8	6.9	119.4
63332_at			AA127696	19.8	11.7	11.4
51155_at			AL043669	19.8	6.2	8.7
45785_at	DKFZP566B1346 protein		AL040982	19.7	8.6	3.4
74908_at	interferon-induced protein 35	IFI35	AW026462	19.7	35.8	8.9
35463_at	homeo box (expressed in ES cells) 1	HESX1	AF059734	19.4	20.3	40.2
91017_at	baculoviral IAP repeat-containing 3	BIRC3	U54711	19.2	19.6	3
48018_at			AA700227	19.1	31.1	9.3
37615_at	growth factor receptor-bound protein 10	GRB10	D86962	19	3.9	60.3
55830_at			AA058686	19	13.5	8.8
38555_at	dual specificity phosphatase 10	DUSP10	AB026436	18.8	7.8	11.2
189_s_at	plasminogen activator, urokinase receptor	PLAUR	U09937	18.8	3.3	5.3
48081_at	myristoylated alanine-rich protein kinase C substrate (MARCKS, 80K-L)	MACS	AA131320	18.7	7.3	8.2
36578_at	baculoviral IAP repeat-containing 2	BIRC2	U37547	18.6	4.9	13.5
73776_at			AI554044	18.6	13.5	14.8
49626_at	Alg5, <i>S. cerevisiae</i> , homolog of		AA425251	18.6	4.7	3.8

Figure 17K

47533_at				AI982765	18.6	12.9	14
32222_at	LR8 protein			AA152202	18.5	14.2	4
44207_at				AI422986	18.5	5.7	5
62561_at	DnaJ (Hsp40) homolog, subfamily B, member 4		DNAJB4	AI741503	18.5	4.6	26.4
37353_g_at	nuclear antigen Sp100		SP100	M60618	18.4	12.6	10
79520_at				AW022213	18.4	8.8	8
32193_at	plexin C1		PLXNC1	AF030339	18.3	38	9.3
36825_at	stimulated trans-acting factor (50 kDa)			X82200	18.3	22.2	11.5
53489_at				AI928764	18.3	8.5	20.8
36918_at	guanylate cyclase 1, soluble, alpha 3		GUCY1A3	Y15723	18.2	3.1	3.6
51510_r_at				T92882	18.1	10.1	4.2
38787_at	promyelocytic leukemia		PML	X63131	17.9	49.4	18.9
53578_at				W70067	17.9	7.9	3.9
1203_at				D17357	17.8	5.2	10.3
34304_s_at	spermidine/spermine N1-acetyltransferase		SAT	AL050290	17.7	12	9.3
42721_at				AI261490	17.5	4.3	5.9
63390_at	KIAA0603 gene product			AI948512	17.5	18.7	12
37823_at	small inducible cytokine subfamily A (Cys-Cys), member 8 (monocyte chemotactic protein 2)		SCYA8	Y16645	17.3	63.9	47.1
35230_at	hypothetical protein, clone 24751			AF070530	17.3	5.7	5.6
50361_at				AI658662	17.1	24.2	11.4
1461_at	nuclear factor of kappa light polypeptide gene enhancer in B-cells inhibitor, alpha		NFKBIA	M69043	17	4.9	6.3
40082_at	fatty acid:Coenzyme A ligase, long-chain 2		FACL2	D10040	16.9	11.5	6.5
39993_at	phosphatidylinositol glycan, class A (paroxysmal nocturnal hemoglobinuria)		PIGA	D11466	16.9	8.5	5.8
53773_at	tumor necrosis factor (ligand) superfamily, member 13b		TNFSF13B	AA682496	16.9	17.6	11.2
54063_at				AI801777	16.9	4.4	6.2
36671_at	asparagine synthetase		ASNS	M27396	16.8	9.8	3.2
33823_at	CD36 antigen (collagen type I receptor, thrombospondin receptor)-like 2 (lysosomal integral membrane protein II)		CD36L2	D12676	16.8	30.9	9.5
55776_at				W20082	16.8	26.9	17.7
56805_at				N52761	16.8	10.4	14.1
40960_at	UDP-Gal:betaGlcNAc beta 1,4-galactosyltransferase, polypeptide 1		B4GALT1	D29805	16.7	4.9	3.9

Figure 17L

67401_at				AI434675	16.7	16.1	6.4
55629_at				AL040635	16.7	9.7	9.2
58569_r_at	niban protein			AI308197	16.7	5.9	4
46655_at	transcription factor (SMIF gene)			AA131327	16.6	8	14.5
39986_at	DKFZP586D0919 protein			AL050100	16.5	4.2	4.9
79721_at				AI733197	16.5	11.6	56.6
52675_at				AI581142	16.5	7.4	3.6
48556_at	HIV TAT specific factor 1		HTATSF1	AA581365	16.4	6.4	4.2
35937_at	MHC class I polypeptide-related sequence B		MICB	U65416	16.3	17.2	32.3
54990_at	hypothetical protein			AW009586	16.3	6	8.2
52062_f_at				H98105	16.3	22.9	6.4
53661_at				AL038450	16.2	5	5.2
72808_at				AI640523	16.1	6.1	32.6
39723_at	cullin 1		CUL1	AF062536	16	6.5	4.8
296_at	tubulin, beta polypeptide		TUBB	X79535	16	12.7	5
41021_s_at	glycerol-3-phosphate dehydrogenase 2 (mitochondrial)		GPD2	U36310	16	10.7	9.1
55673_at	RAB-8b protein			AI807023	16	13.1	6.6
36600_at	proteasome (prosome, macropain) activator subunit 1 (PA28 alpha)		PSME1	L07633	15.9	19.1	6.7
35149_at	tumor necrosis factor receptor superfamily, member 5		TNFRSF5	AI865431	15.8	11.1	6.1
82359_at				AI347618	15.7	26.5	44.6
40088_at	nuclear receptor interacting protein 1		NRIP1	X84373	15.6	9.9	8.5
60016_at				AI400414	15.6	9	9.5
86261_at				AA017070	15.5	12.3	7.8
38006_at	CD48 antigen (B-cell membrane protein)		CD48	M37766	15.4	21.6	9.8
40755_at	MHC class I polypeptide-related sequence A		MICA	X92841	15.4	8	13.9
69589_at				AI743005	15.4	14.3	42.7
56278_at				AI692445	15.4	6.3	58
42933_at				T95101	15.3	4.8	11.4
1052_s_at	CCAAT/enhancer binding protein (C/EBP), delta		CEBPB	M83667	15.2	3	7
38287_at	proteasome (prosome, macropain) subunit, beta type, 9 (large multifunctional protease 2)		PSMB9	AA808961	15.2	15.8	3
37732_at	RING1 and YY1 binding protein		RYBP	AL049940	15.1	4.6	5.3
81726_at				AI885498	15.1	10.7	7.2
63984_f_at				AA535914	15	16.4	9.2

Figure 17M

48039_at				HLA-A	AI634580	15	16.9	9.4
41237_at	major histocompatibility complex, class I, A				D32129	14.9	9.7	5.6
87002_i_at					AI270476	14.8	28.7	63.7
40556_at	reticulocalbin 1, EF-hand calcium binding domain			RCN1	D42073	14.7	9.7	5.4
55051_at					AA005281	14.7	16.7	12.4
51672_f_at					AI161358	14.7	5.7	16.9
40957_at	KIAA0160 protein			KIAA0160	D63881	14.6	17.7	5
33804_at	protein tyrosine kinase 2 beta			PTK2B	U43522	14.6	13.4	5
32533_s_at	vesicle-associated membrane protein 5 (myobrevin)			VAMP5	AF054825	14.5	39.7	13.4
78077_at					AI125204	14.5	6	3
51854_at	KIAA1128 protein				AA424126	14.4	6.3	5.6
31680_at					M55630	14.3	5.5	6.7
38649_at	KIAA0970 protein				AB023187	14.2	7	30.1
36992_at	thioredoxin			TXN	AI653621	14.2	9.8	5.8
56889_at	dual specificity phosphatase 10			DUSP10	N36770	14.2	4.1	10.6
70516_at	leucine-rich protein mRNA				AI817736	14.2	6.4	8.6
46526_at	hypothetical protein MGC5242				AI434025	14.2	13.4	13.9
59516_at	KIAA1268 protein				AA210695	14.1	11.3	10.5
34308_at	H2A histone family, member L			H2AFL	U90551	14	4.9	13.2
54064_i_at					AA903287	14	5	5.7
51798_at	hypothetical protein FLJ11000				AW006887	14	13	3.9
36872_at	cyclic AMP phosphoprotein, 19 kD				AL120559	13.9	12.3	4.7
49302_at	HBV associated factor				AI971817	13.9	9.1	5.4
48472_at	NY-REN.18 antigen				AI425002	13.9	8.8	3.9
41532_at	zinc finger protein 151 (pHZ-67)			ZNF151	Y09723	13.8	5.4	4.2
62314_at					H04394	13.8	3.7	4.6
56832_at					AI871408	13.7	6.9	12.3
74566_at	insulin-like growth factor-binding protein 4			IGFBP4	AI440424	13.7	75.4	14.1
51508_i_at					T92882	13.7	7.2	9.7
32069_at	KIAA0615 gene product				AB014515	13.5	7.4	10
52167_at					AA151346	13.5	12.1	11.7
38207_at					AW006742	13.4	6.4	12.4
37360_at	lymphocyte antigen 6 complex, locus E			LY6E	U66711	13.4	13.6	8.3
61118_at					AI983632	13.4	8.3	3.1
42362_at	CCAAT/enhancer binding protein (C/EBP), delta			CEBPD	AI633245	13.4	3.1	3.4

Figure 17N

41474_at	kinesin heavy chain member 2	KIF2	Y08319	13.3	24.6	7.3
39708_at	signal transducer and activator of transcription 3 (acute-phase response factor)	STAT3	L29277	13.3	5.3	4.4
51024_at			AI400509	13.3	17.3	7.2
36776_at	lymphocyte-activation gene 3	LAG3	X51985	13.2	148.7	69.7
33106_at	nuclear receptor subfamily 1, group H, member 3	NR1H3	U22662	13.2	18.1	18
1170_at	colony stimulating factor 1 (macrophage)	CSF1	M37435	13.1	6.9	3.2
843_at	protein tyrosine phosphatase type IVA, member 1	PTP4A1	U48296	13.1	3.7	7.2
36103_at	small inducible cytokine A3 (homologous to mouse Mip-1a)	SCYA3	D90144	13.1	6.1	7.6
51220_at	small EDRK-rich factor 1A (telomeric)	SERF1A	W67895	13.1	4.7	3.6
56080_at			AI703473	13.1	12.8	8.3
44838_at	hypothetical protein FLJ12752		AI271348	13	4	3.1
40137_at	protein tyrosine phosphatase, non-receptor type 1	PTPN1	M31724	12.9	5.3	5.2
36484_at	UDP-N-acetyl-alpha-D-galactosamine:polypeptide N-acetylgalactosaminyltransferase 3 (GalNAc-T3)	GALNT3	AI935146	12.9	5.5	8.4
61489_at			AI718763	12.9	3.7	3.4
44853_s_at	CGI-34 protein		AA528059	12.9	10	7.4
39628_at	RAB9, member RAS oncogene family	RAB9	AI671547	12.8	7.4	3.5
39857_at	syntaxin 11	STX11	AF044309	12.7	5.6	12.9
44920_at			AI914925	12.7	4.8	7.6
32217_at			AF052105	12.6	6.3	7.6
37049_g_at	translocase of outer mitochondrial membrane 34		U58970	12.6	17.9	13.6
45721_at	endothelin converting enzyme 1	ECE1	AL039866	12.6	4.3	5.7
63662_at			AI126468	12.6	4.9	4.5
33886_at	spectrin SH3 domain binding protein 1	SSH3BP1	AF006516	12.4	4.4	6.4
82094_i_at			AI810266	12.4	6.4	36.3
56804_at			AA916789	12.3	14.3	3.6
55400_at	major histocompatibility complex, class I, F	HLA-F	AI732773	12.3	4.2	5.8
57699_at			AA133285	12.1	31.2	65.3
34446_at	KIAA0471 gene product		AL049701	12	39.6	24.8
1909_at	B-cell CLL/lymphoma 2	BCL2	M14745	12	9.9	8.1
38439_at	nuclear factor (erythroid-derived 2)-like 1	NFE2L1	L24123	12	7.5	3.5
60017_at			AA142842	12	11.9	5.7
45296_at	KIAA1631 protein		AI986035	12	13	4.3

Figure 170

50320_g_at				AI497833	12	3.8	5.7
61940_at	hypothetical protein FLJ21343			W28128	12	28.9	20.7
1761_at	platelet-derived growth factor receptor-like	PDGFR		D37965	11.9	34.7	14.4
72862_at				AW026509	11.9	15.3	10
39821_s_at	growth arrest and DNA-damage-inducible, beta	GADD45B		N95168	11.8	7.1	21.4
45608_at				AI202327	11.7	4.7	7.2
195_s_at	caspase 4, apoptosis-related cysteine protease	CASP4		U28014	11.6	8.7	4.5
64001_s_at	seven in absentia (Drosophila) homolog 2	SIAH2		AW008929	11.6	4.2	5.3
1842_at				S62138	11.6	25	46.1
39420_at				S62138	11.6	20.2	42.5
32776_at	v-rat simian leukemia viral oncogene homolog B (ras related; GTP binding protein)	RALB		M35416	11.3	5.2	6.4
1358_s_at	interferon, alpha-inducible protein (clone IFI-6-16)	G1P3		U22970	11.3	8.6	10
49943_at				AI681558	11.3	3.8	8.3
54933_at				AI356461	11.3	8.8	4.7
55001_at				R13843	11.2	15.2	9.2
1089_i_at				M64936	11.1	3.3	13.9
64174_at				AI127460	11	8.6	7
85409_at	neuropilin 2	NRP2		AI253672	10.9	11.1	12.7
41304_at	dual specificity phosphatase 4	DUSP4		AF016098	10.8	5.7	5.9
1788_s_at	butyrophilin, subfamily 3, member A1	BTN3A1		U48807	10.8	35.6	12.4
32629_f_at	protein kinase, AMP-activated, beta 1 non-catalytic subunit	PRKAB1		U90552	10.8	9.9	8.4
35348_at				AF022116	10.8	3	5.9
42365_f_at				H60126	10.8	6.9	3.3
64692_at				W30810	10.8	3.6	16.2
41690_at				AL049471	10.7	3.2	5.4
36456_at	DKFZP564I052 protein			AL080063	10.7	6.9	4.7
37536_at	CD83 antigen (activated B lymphocytes, immunoglobulin superfamily)	CD83		Z11697	10.7	8.3	6.3
44742_s_at	hypothetical protein FLJ21939 similar to 5-azacytidine induced gene 2			AA205994	10.7	5.1	4.7
59478_at				AA741304	10.6	26.4	11.9
55551_at				AI766561	10.6	7.1	10.3
43750_at	tumor necrosis factor receptor superfamily, member 10a	TNFRSF10A		AA102746	10.6	4.2	6.2

Figure 17P

53139_at	protein phosphatase 1B (formerly 2C), magnesium-dependent, beta isoform	PPM1B	AW020971	10.6	7.4	11.9
34749_at	solute carrier family 31 (copper transporters), member 2	SLC31A2	U83461	10.5	12.9	5.5
37754_at	lectin, galactoside-binding, soluble, 3 binding protein (galectin 6 binding protein)	LGALS3BP	L13210	10.5	19.8	7.3
52226_at			H16779	10.5	3.6	4.2
50427_at			AW003857	10.5	7.2	3.5
55809_at			AA028139	10.5	3.3	3.8
1402_at	v-yes-1 Yamaguchi sarcoma viral related oncogene homolog	LYN	M16038	10.4	7.2	5.2
48080_at	hypothetical protein FLJ20037		AI566481	10.4	12.6	45.7
67087_at			AI351909	10.3	4.9	4.9
68047_at			AA936632	10.3	14	5.6
64176_at			AI650542	10.3	6.9	8.5
39703_at	ubiquitously transcribed tetratricopeptide repeat gene, X chromosome	UTX	AF000993	10.2	4.6	48.9
57051_at			AA167704	10.2	9.4	6.1
86737_at			H88330	10.2	6	4.1
45261_at	leucine aminopeptidase		AF034175	10.1	15	8.1
50297_f_at	apolipoprotein L, 3	APOL3		10	17.1	7.3
38517_at	interferon-stimulated transcription factor 3, gamma (48kD)	ISGF3G	M87503	10	6.3	11.4
36294_at	serine/threonine kinase 4	STK4	U60207	10	10.6	4.4
46079_at			W63785	10	6	5.1
34740_at	forkhead box O3A	FOXO3A	AF032886	9.9	3.5	6.3
33847_s_at	cyclin-dependent kinase inhibitor 1B (p27, Kip1)	CDKN1B	AI304854	9.9	5	8.2
37674_at	aminolevulinatase, delta-, synthase 1	ALAS1	Y00451	9.9	3.7	12.5
70151_at			AI805522	9.8	6.3	5.3
49650_at			N68715	9.8	7.4	3.1
40146_at	RAP1B, member of RAS oncogene family	RAP1B	AL080212	9.7	4.3	5.7
32088_at	basic leucine zipper nuclear factor 1 (JEM-1)	BLZF1	U79751	9.7	6.4	16.3
1478_at	IL2-inducible T-cell kinase	ITK	L10717	9.7	4	7
37218_at	BTG family, member 3	BTG3	D64110	9.7	8.7	6.5
51131_at			AW007736	9.7	8.3	4.6
62512_at			AI927692	9.7	7.3	18.4
49080_at			AL042923	9.7	5.8	9.9

Figure 17Q

1061_at	interleukin 10 receptor, alpha	IL10RA	U00672	9.6	5.1	11.1
57537_at			AA001250	9.6	8.5	4.6
54857_at	ancient conserved domain protein 4		AI636759	9.6	8	7.1
43463_at	neuropilin 2	NRP2	N90777	9.6	7.2	11.4
53498_at			AA464480	9.6	7	6.1
66326_at			AI936888	9.5	5	3.8
42720_at			AI393727	9.5	3.2	11.3
43225_i_at			AA001052	9.5	4	4.8
44673_at			N53555	9.5	13.7	4.3
39976_at	hypothetical protein from EUROIMAGE 783647		AL050087	9.4	4.1	5.9
65795_r_at			AL042362	9.4	3	4.7
35310_at			D45288	9.4	5.5	8.9
33598_r_at	chromosome 1 open reading frame 7	C1ORF7	AF054176	9.3	3.4	3.2
1184_at	proteasome (prosome, macropain) activator subunit 2 (PA28 beta)	PSME2	D45248	9.3	10.3	5.7
44763_at			AA280886	9.3	3.8	6.9
33227_at	interleukin 10 receptor, beta	IL10RB	AI984234	9.2	4.2	3.1
33236_at	retinoic acid receptor responder (tazarotene induced) 3	RARRES3	AF060228	9.2	55.2	24.8
1323_at	ubiquitin B	UBB	X04803	9.2	4.8	3.4
287_at	activating transcription factor 3	ATF3	L19871	9.2	4.8	15.2
53720_at	hypothetical protein FLJ11286		AI862559	9.2	10.4	10.1
45258_at	hypothetical protein DKFp761G2113		AF039240	9.2	4.7	3.2
38149_at	KIAA0053 gene product		D29642	9.1	8.7	4.3
41156_g_at	catenin (cadherin-associated protein), alpha 1 (102kD)	CTNNA1	U03100	9.1	5.3	3.7
72290_r_at			AA814901	9.1	10.9	12.8
52535_s_at	cyclin-dependent kinase inhibitor 1B (p27, Kip1)	CDKN1B	AI680672	9.1	3.9	8.3
49508_at			AI041540	9.1	5.2	3.5
71661_at			AI174651	9.1	5.4	6.5
61675_at			AI949662	9.1	12.7	14.2
43848_s_at			AI968465	9.1	8.8	27.3
78605_at			AI080388	9	4.4	5.6
41830_at	KIAA0494 gene product		AB007963	9	5.6	6.4
41164_at	immunoglobulin heavy constant mu	IghM	X67301	8.9	8.3	47.2
37544_at	nuclear factor, interleukin 3 regulated	NFIL3	X64318	8.9	8.7	3.2
37603_at	interleukin 1 receptor antagonist	IL1RN	X52015	8.9	5.9	5.6

Figure 17R

37915_at	bromodomain adjacent to zinc finger domain, 2B	BAZ2B	AL080173	8.9	3.4	3.9
45233_at	mesenchymal stem cell protein DSC92		W22743	8.9	4.4	3.6
44561_r_at			AI096866	8.9	9.6	6.4
44851_s_at			N48088	8.8	8.4	3.5
51121_at			R45590	8.8	4.6	5.2
64739_at	docking protein 2, 56kD	DOK2	AI828929	8.8	8.3	8.5
37126_at	Sjogren syndrome antigen A1 (52kD, ribonucleoprotein autoantigen SS-A/Ro)	SSA1	M62800	8.6	8.2	10.6
35337_at	F-box only protein 7	FBXO7	AL050254	8.6	8.5	16.3
54875_at			N57875	8.6	4.9	12
64457_at	leucine rich repeat (in FLII) interacting protein 2	LRRFIP2	AA203144	8.6	4.4	3.8
46858_at			AI796221	8.5	8.5	4.4
41169_at	plasminogen activator, urokinase receptor	PLAUR	X74039	8.4	4.2	4
91669_at			AA491645	8.4	6.2	5.6
37684_at	solute carrier family 21 (organic anion transporter), member 9	SLC21A9	AB020687	8.4	6.8	8
612_s_at	2',3'-cyclic nucleotide 3' phosphodiesterase	CNP	M19650	8.3	3.9	5.2
40468_at	KIAA0554 protein		AB011126	8.3	9	5.1
1449_at	proteasome (prosome, macropain) subunit, alpha type, 4	PSMA4	D00763	8.3	6.2	10.1
83378_at			T54678	8.3	3.3	6.3
89412_at			AI804386	8.3	3.5	5.3
57543_at	bromodomain adjacent to zinc finger domain, 1A	BAZ1A	AA112166	8.3	6.2	6.5
35136_at	hypothetical protein P15-2		AL031387	8.2	4.5	9.7
41488_at	hypothetical protein A-211C6.1		AC002394	8.2	4.9	4
276_at	DnaJ (Hsp40) homolog, subfamily A, member 1	DNAJA1	L08069	8.2	5.2	6
44782_at			AA258063	8.2	4.3	5.9
33352_at	H2A histone family, member Q	H2AFQ	X57985	8.1	11.8	37.7
36753_at	leukocyte immunoglobulin-like receptor, subfamily B (with TM and ITIM domains), member 4	LILRB4	AF072099	8.1	8.9	3.9
36943_r_at	pleiomorphic adenoma gene-like 1	PLAGL1	U81992	8.1	5.8	7.6
42513_at	ATPase, Na+/K+ transporting, beta 3 polypeptide	ATP1B3	AA151771	8.1	4.7	6.4
32218_at			AF034176	8.1	8.1	5.5
33859_at	sin3-associated polypeptide, 18kD	SAP18	U96915	8	12.8	10.9
61757_at			AI273692	8	3.3	7.4
63394_at	RBP1-like protein		AI365221	8	4.2	9

Figure 17S

40063_at	nuclear domain 10 protein			U22897	7.9	5.1	9.3
38986_at	glucose regulated protein, 58kD	GRP58		Z49835	7.9	16.1	4.7
37363_at	KIAA0429 gene product			AB007889	7.9	7.6	3.8
55194_at				AI638607	7.9	4.1	4.8
32128_at	small inducible cytokine subfamily A (Cys-Cys), member 18, pulmonary and activation-regulated	SCYA18		Y13710	7.8	7.9	4
36472_at	N-myc (and STAT) interactor	NMI		U32849	7.8	12.1	8.6
35321_at	tousled-like kinase 2	TLK2		AB004884	7.8	7.3	4.5
36934_at	hypothetical protein LOC57158			AL035447	7.8	3.9	13.6
32595_at	G-rich RNA sequence binding factor 1	GRSF1		U07231	7.8	22.7	16.5
38014_at	adenosine deaminase, RNA-specific	ADAR		X79448	7.8	7.4	5.9
56401_at	zinc finger protein 22 (KOX 15)	ZNF22		AI951311	7.8	4.7	6.3
57675_at				AW052119	7.8	5.1	3.3
45234_at				AA648522	7.8	11.3	14.9
54826_at				AA284268	7.8	4.8	3.3
67470_at				AI928355	7.8	4.5	6.4
45165_at				AI831506	7.8	3.2	4.8
44499_at				N67390	7.8	3.8	4.6
45055_at				AA461205	7.8	5	5.5
32065_at	cAMP responsive element modulator	CREM		S68134	7.7	8.5	47.4
33799_at	seven in absentia (Drosophila) homolog 2	SIAH2		U76248	7.7	3.6	5.4
36122_at	proteasome (prosome, macropain) subunit, alpha type, 6	PSMA6		X59417	7.7	3.3	4.7
83245_at				AI300111	7.7	4.9	9.2
48566_at	hypothetical protein FLJ20446			N98615	7.7	6.4	3.8
50266_at				AI741178	7.7	6.5	5.4
74689_at	lymphocyte-activation gene 3	LAG3		AW003135	7.7	77.5	26.1
44688_at	RelA-associated inhibitor			AA614049	7.7	6.7	8.8
41329_at	hypothetical protein LOC57147			AI458463	7.6	8.1	5.7
38676_at	stress 70 protein chaperone, microsome-associated, 60kD	STCH		AA059408	7.6	3.8	4
237_s_at	protein phosphatase 2 (formerly 2A), catalytic subunit, alpha isoform	PPP2CA		M60483	7.6	5	8.6
57501_at				AI914124	7.6	12.8	5.9
43269_at	KIAA1554 protein			AA424170	7.6	12.5	10.4
75024_at	adenosine deaminase, RNA-specific	ADAR		R49062	7.6	9.5	4

Figure 17T

58691_at				AI420336	7.6	7	4.5
32227_at	proteoglycan 1, secretory granule	PRG1		X17042	7.5	5.4	4.5
51183_at	transcriptional intermediary factor 1 gamma			N98658	7.5	4.9	4.9
49903_at	KIAA1404 protein			W72798	7.5	3.2	6.5
89128_at	C-type (calcium dependent, carbohydrate-recognition domain) lectin, superfamily member 9	CLECSF9		AI968491	7.5	4.9	6.6
68299_at				R07844	7.5	5.6	4.4
45893_at				H24387	7.5	5	7.6
38114_at	RAD21 (S. pombe) homolog	RAD21		D38551	7.4	5.2	5.1
58411_at	KIAA1610 protein			AL037424	7.4	5.4	12.2
32842_at	B-cell CLL/lymphoma 7A	BCL7A		X89984	7.4	5.2	6.1
64508_s_at	KIAA0430 gene product			AI968969	7.4	6.4	8.5
61865_at				AI423165	7.3	4	3.9
76098_r_at				AI304317	7.3	10	6
48742_at	activated RNA polymerase II transcription cofactor 4			AI979012	7.3	7.4	3.6
89856_at	CD83 antigen (activated B lymphocytes, immunoglobulin superfamily)	CD83		AI983994	7.3	7.3	5.3
1326_at	caspase 10, apoptosis-related cysteine protease	CASP10		U60519	7.2	18.6	15.9
36845_at	KIAA0136 protein			D50926	7.2	5.1	5.4
59198_at	hypothetical protein			AF075050	7.2	4.3	3.6
43697_at				AA504249	7.2	3.3	3.4
43553_at				AI807404	7.2	6.3	3.1
40585_at	adenylate cyclase 7	ADCY7		D25538	7.2	5.1	7.2
65827_at	STIP1 homology and U-Box containing protein 1	STUB1		AI630895	7.2	6.3	8.1
40639_at	SCO (cytochrome oxidase deficient, yeast) homolog 2	SCO2		AL021683	7.1	5.2	30.4
39132_at	SWI/SNF related, matrix associated, actin dependent regulator of chromatin, subfamily a, member 5	SMARCA5		AB010882	7.1	3.9	16.4
62431_at				AI197909	7.1	7.9	7.6
56697_r_at				AI791520	7.1	8.2	8.3
1046_at	retinoic acid- and interferon-inducible protein (58kD)			U34605	7	5.4	3
73703_at				AA777616	7	4.7	7.5
59008_i_at				AI815758	7	4.2	12.5
63683_at				N56920	7	7	11
64277_i_at	hypothetical protein FLJ20647			AI478759	7	14.4	9.8
46183_at				AI810826	7	6.2	4.6

Figure 17U

79341_at				AA084622	7	12	17.6
33380_at	topoisomerase-related function protein 4.1			AB005754	6.9	6.7	20
35738_at	high-mobility group (nonhistone chromosomal) protein 17-like 3	HMG17L3		AI347088	6.9	3.9	3.6
37026_at	core promoter element binding protein	COPEB		AF001461	6.9	5.7	8.6
35302_at	nuclear RNA export factor 1	NXF1		AJ132712	6.9	8.3	34.6
34041_at	small inducible cytokine subfamily A (Cys-Cys), member 22	SCYA22		U83171	6.9	5	5.4
65135_at				AI916853	6.9	3.6	4.9
67223_at				AI003757	6.9	7.8	6.5
59344_at				AA017033	6.9	3	4.8
63083_r_at				R63071	6.9	3.4	5.8
57785_at	H4 histone family, member H	H4FH		N32748	6.9	5.8	9.3
46961_r_at				T91186	6.9	8.3	9.6
44120_at				AI879381	6.9	6.9	7.4
32203_at	HBV associated factor			AA160708	6.8	8.2	5.5
37391_at	cathepsin L	CTSL		X12451	6.8	9.3	4.2
62246_at				T92947	6.8	3.4	20.8
79735_at				AA037529	6.8	3.7	3.9
38712_at	membrane protein CH1			AL035291	6.7	6.1	6.9
41235_at	activating transcription factor 4 (tax-responsive enhancer element B67)	ATF4		AL022312	6.7	4.1	3.1
37971_at	bromodomain adjacent to zinc finger domain, 1A	BAZ1A		AL050089	6.7	6.9	6.4
39158_at	activating transcription factor 5	ATF5		AB021663	6.7	6.6	4.7
56435_at	hypothetical protein FLJ20625			AA573749	6.7	5.9	8.5
39710_at	P311 protein			U30521	6.6	35	8.7
40803_at				AL050161	6.6	3.6	4.5
32725_at	BH3 interacting domain death agonist	BID		AF042083	6.6	4.2	9.3
32784_at	serine/threonine-protein kinase PRP4 homolog			AB011108	6.6	3.2	3.3
59000_at	x 010 protein	x 010		AA629304	6.6	4.8	4
44718_at				AA577672	6.6	5.7	7.9
53666_at				AA114900	6.6	3.8	6.4
33777_at	thromboxane A synthase 1 (platelet, cytochrome P450, subfamily V)	TBXAS1		D34625	6.6	6.5	6.3

Figure 17V

37275_at	GA-binding protein transcription factor, beta subunit 1 (53kD)	GABPB1	U13045	6.5	3.9	10.2
48929_at			AA521311	6.5	10	3.7
74456_r_at			AI970474	6.5	5.5	4
32563_at	ATPase, Na+/K+ transporting, beta 3 polypeptide	ATP1B3	U51478	6.4	3.6	3.8
45379_at			AI659019	6.4	5.8	9.6
76364_at			AA643507	6.4	17.1	23.3
43387_s_at	hypothetical protein FLJ13171		AW001604	6.4	13.8	9.9
47655_at			AI022328	6.4	6.5	8.6
1385_at	transforming growth factor, beta-induced, 68kD	TGFB1	M77349	6.4	3.5	6.3
58954_at	NADH dehydrogenase (ubiquinone) Fe-S protein 8 (23kD) (NADH-coenzyme Q reductase)	NDUFS8	AW021179	6.4	5.3	6.5
34951_at	putative chemokine receptor; GTP-binding protein		D10923	6.3	3.2	4.6
49875_at			AI733353	6.3	5.9	16.6
65364_at			W56309	6.3	9.2	8.1
53128_at	hypothetical protein FLJ10494		AI678076	6.3	5	3.2
44080_at			W68180	6.3	5.2	5.5
35328_at			AF055023	6.2	3.4	7.9
38805_at	TG-interacting factor (TALE family homeobox)	TGIF	X89750	6.2	4.4	6.6
63013_at			AL120446	6.2	7.7	4.2
55525_at			AI972498	6.2	5.5	6.3
40570_at	forkhead box O1A (rhabdomyosarcoma)	FOXO1A	AF032885	6.1	8.1	4.5
566_at	promyelocytic leukemia	PML	M79462	6.1	6.6	8.4
55239_at			AA233120	6.1	7.4	6.8
53708_at	cation-chloride cotransporter-interacting protein		AI467928	6.1	5.6	5.2
53881_s_at			AL044098	6.1	5.1	5.8
34178_at	zinc finger protein 297	ZNF297	AI884738	6	3.4	5.3
33360_at	f-box and leucine-rich repeat protein 11	FBXL11	AB023221	6	3.5	4.9
38662_at			AL047596	6	9.1	4.5
37221_at	protein kinase, cAMP-dependent, regulatory, type II, beta	PRKAR2B	M31158	6	4.6	5
58192_at	KIAA1411 protein		AI308839	6	4.9	3.7
47513_at			AA679424	6	4.2	5.3
79858_at			AI693951	6	5.7	3.2
48431_at	fucosyltransferase 4 (alpha (1,3) fucosyltransferase, myeloid-specific)	FUT4	AI970292	6	14.7	13.8

Figure 17W

35789_at	KIAA1042 protein		AB028965	6	6	6.4
33499_s_at	immunoglobulin heavy constant gamma 3 (G3m marker)	IGHG3	AF067420	5.9	3.8	17.1
894_g_at	ubiquitin carrier protein		M91670	5.9	7.8	5.4
41183_at	cleavage stimulation factor, 3' pre-RNA, subunit 3, 77kD	CSTF3	U15782	5.9	4.6	6.3
35936_g_at	carnitine palmitoyltransferase I, muscle	CPT1B	Y08683	5.9	18.4	6.7
51643_at			AL036254	5.9	3.9	6.5
65351_r_at			AA146979	5.9	3.2	5.1
91341_at			AI291659	5.9	4.5	3.8
41295_at	GTT1 protein		AL041780	5.9	2.9	7.4
32550_r_at	CCAAT/enhancer binding protein (C/EBP), alpha	CEBPA	Y11525	5.9	4.9	4.6
49003_at	NADH dehydrogenase (ubiquinone) 1 beta subcomplex, 10 (22kD, PDSW)	NDUFB10	AA195976	5.9	5.8	5.9
37424_at	HCR (a-helix coiled-coil rod homologue)		AB029343	5.8	3.2	3.5
36535_at	microfibrillar-associated protein 1	MFAP1	U04209	5.8	5.1	7.1
50713_at			H73401	5.8	4.1	6.6
82594_at			AI308862	5.8	6.1	3.1
50210_at			AA429326	5.8	19.9	21.3
38828_s_at	KH-type splicing regulatory protein (FUSE binding protein 2)	KHSRP	AA628946	5.8	2.8	5.8
54152_at	eukaryotic translation initiation factor 4E binding protein 1	EIF4EBP1	AI026669	5.8	5.2	5.8
41126_at	solute carrier family 1 (glutamate/neutral amino acid transporter), member 4	SLC1A4	AA978353	5.7	7.9	4.7
39136_at	oxidative-stress responsive 1	OSR1	AB017642	5.7	3.1	3
59314_at	retinoic acid induced 1	RAI1	AI926671	5.7	4.5	3.2
47144_at	ASB-1 protein		AI018173	5.7	4.1	4.5
1357_at	ubiquitin specific protease 4 (proto-oncogene)	USP4	U20657	5.7	3.4	8.3
46664_at	apoptosis-associated speck-like protein containing a CARD		AI148558	5.7	5.1	6.3
40791_at	polymerase (RNA) II (DNA directed) polypeptide A (220kD)	POLR2A	X63564	5.6	3.1	4.2
33418_at	RAB3 GTPase-ACTIVATING PROTEIN		AL096752	5.6	7.2	5.2
35251_at	human immunodeficiency virus type I enhancer-binding protein 1	HIVEP1	X51435	5.6	6.4	12.2
1295_at	v-rel avian reticuloendotheliosis viral oncogene homolog A (nuclear factor of kappa light polypeptide gene enhancer in B-cells 3 (p65))	RELA	L19067	5.6	3.8	5.3

Figure 17X

71936_at				AI148006	5.6	4.2	3.2
51185_at	PAN2 protein			AI309334	5.6	4.2	3.3
53302_r_at				AI741209	5.6	4.3	10.1
40045_g_at	chromosome 18 open reading frame 1		C18ORF1	AF009425	5.6	4.5	3.1
49390_at				AA999858	5.6	5.6	5.6
52261_at				AI970054	5.6	4.7	6.4
41432_at	ribosomal protein S6 kinase, 90kD, polypeptide 5		RPS6KA5	AF090421	5.5	3.3	8.7
39061_at	bone marrow stromal cell antigen 2		BST2	D28137	5.5	9.7	6.6
1310_at	proteasome (prosome, macropain) subunit, beta type, 2		PSMB2	D26599	5.5	7	4.6
41355_at	E2F transcription factor 5, p130-binding		E2F5	N95229	5.5	5.7	8
409_at	tyrosine 3-monooxygenase/tryptophan 5-monooxygenase activation protein, theta polypeptide		YWHAQ	X56468	5.5	3.3	6.2
74975_f_at	zinc finger protein 173		ZNF173	AW009746	5.5	5.5	14.1
75201_f_at				AI688900	5.5	11	7
65814_at				N95618	5.5	5.2	4.5
68328_at				AI475514	5.5	10.2	32.1
35247_at	small nuclear RNA activating complex, polypeptide 5, 19kD			AI557062	5.5	2.9	5
36991_at	splicing factor, arginine/serine-rich 4		SFRS4	L14076	5.4	5.3	4.2
91400_at				AI309016	5.4	3.6	3.2
63526_f_at				AI669667	5.4	5.8	7.1
45521_s_at	KIAA0100 gene product			AI967955	5.4	5.5	5.7
63793_at	DNA polymerase epsilon p12 subunit			AI797479	5.4	3.7	4.8
45736_at	two-pore channel 1, homolog			AL048491	5.4	5.2	5.1
1383_at	protein phosphatase 2 (formerly 2A), regulatory subunit B (PR 52), alpha isoform		PPP2R2A	M64929	5.3	7.5	12.4
38778_at	KIAA1046 protein			AB028969	5.3	4.8	3.3
61840_s_at	catenin (cadherin-associated protein), delta 1		CTNND1	AI863000	5.3	4.5	9.9
59451_at				AI800640	5.3	6.2	23.2
46744_s_at	carboxypeptidase M		CPM	AA902480	5.3	3.6	3.5
85691_at				AI989354	5.3	5.6	6.1
70228_r_at	protein kinase, cAMP-dependent, regulatory, type II, beta		PRKAR2B	AI792179	5.3	9.7	5
67220_at				AW005511	5.3	3	3
40409_at	aldehyde dehydrogenase 10 (fatty aldehyde dehydrogenase)		ALDH10	U46689	5.3	3.9	5.1

Figure 17Y

40822_at	nuclear factor of activated T-cells, cytoplasmic, calcineurin-dependent 3	NFATC3	L41067	5.3	4.4	3.8
35336_at	KIAA0063 gene product		AL021707	5.3	3.6	6.2
59721_at	clone FLB4739		AI973147	5.3	4.8	6.4
55567_at			AA147933	5.3	5.3	5.2
39742_at	TRAF family member-associated NFKB activator	TANK	U59863	5.2	4	3.1
33869_at			AL080218	5.2	3.2	3.8
277_at	myeloid cell leukemia sequence 1 (BCL2-related)	MCL1	L08246	5.2	3.9	10
48873_at	Gene 33/Mig-6		AI571452	5.2	3.4	25.2
80829_at			AA210905	5.2	4	3.4
50205_at	leptin receptor	LEPR	N46838	5.2	5.1	15
45624_at	core1 UDP-galactose:N-acetylgalactosamine-alpha-R beta 1,3-galactosyltransferase		AI967968	5.2	6.9	4
55915_r_at	hypothetical protein LOC54149		AI922520	5.2	13.7	15.9
40955_at	prostate tumor over expressed gene 1	PTOV1	U79287	5.2	2.6	4.6
41540_at	protein phosphatase 1, regulatory subunit 7	PPP1R7	Z50749	5.2	2.9	5.2
34811_at	ATP synthase, H+ transporting, mitochondrial F0 complex, subunit c (subunit 9) isoform 3	ATP5G3	U09813	5.2	3.2	6.2
36601_at	vinculin	VCL	M33308	5.2	4.6	5.1
37002_at	biliverdin reductase B (flavin reductase (NADPH))	BLVRB	D32143	5.2	3.8	5.3
48750_at	class I cytokine receptor		AI983115	5.2	4.6	4.9
44638_at	insulin-degrading enzyme	IDE	AA573292	5.2	2.6	4.2
59136_at			AA779895	5.2	5.1	5.2
48811_at	N-acetylglucosamine-1-phosphodiester alpha-N-acetylglucosaminidase		AW007594	5.2	4	5
51123_at			H11724	5.2	5	4.9
53120_at	hypothetical protein FLJ20442		AI681916	5.2	4.3	4.8
32792_at	GCLP-interacting protein p29		AL031432	5.1	3.2	4.1
38369_at	myeloid differentiation primary response gene (88)	MYD88	U70451	5.1	15	18.4
38824_at	Tat-interacting protein (30kD)		AF039103	5.1	5.8	12.6
84981_r_at	collagen, type III, alpha 1 (Ehlers-Danlos syndrome type IV, autosomal dominant)	COL3A1	T54146	5.1	8.6	5.6
50763_at	KIAA1080 protein; Golgi-associated, gamma-adaptin ear containing, ARF-binding protein 2		AA779863	5.1	6	3.4
44397_at			H11252	5.1	4.3	4.4

Figure 17Z

45181_at	zinc finger protein 281	ZNF281	AA121673	5.1	5.9	9.5
1630_s_at	spleen tyrosine kinase	SYK	Z29630	5.1	3.9	4.9
33833_at	spectrin, alpha, non-erythrocytic 1 (alpha-fodrin)	SPTAN1	J05243	5.1	3.8	5.7
61853_at			AA773816	5.1	5.1	5.1
45224_at			AA025862	5.1	3.2	3.5
55695_at	glycine cleavage system protein H (aminomethyl carrier)	GCSH	AI890917	5.1	4.5	4.2
32142_at	serine/threonine kinase 3 (Ste20, yeast homolog)	STE3	U26424	5	4.1	3.6
32233_at	torsin family 1, member B (torsin B)	TOR1B	AF007872	5	6.9	13
49516_at			AA531016	5	6.6	3.5
36270_at	CD86 antigen (CD28 antigen ligand 2, B7-2 antigen)	CD86	U04343	4.9	13.2	3.9
41592_at	JAK binding protein		AB000734	4.9	6	7.7
45612_at			AI744560	4.9	3.7	5.9
55558_at			AA524361	4.9	4.8	3.6
34406_at	KIAA0602 protein		AB011174	4.9	2.7	4.6
31722_at	ribosomal protein L3	RPL3	AL022326	4.9	3.9	9
56546_at	KIAA1545 protein		H17730	4.9	5.9	5.8
53884_at			AA312905	4.9	3.9	4.7
43053_g_at			AW025012	4.9	3.2	4.6
51239_at	hypothetical protein FLJ23239		W67828	4.9	3.8	4.4
49633_at	hypothetical protein HES6		N21131	4.9	4.9	4.8
54878_at			AL048421	4.9	3.8	4.8
41572_r_at	v-rel avian reticuloendotheliosis viral oncogene homolog	REL	X75042	4.8	3.5	3.7
39182_at	epithelial membrane protein 3	EMP3	U87947	4.8	3.8	5
50829_at			AA706818	4.8	10.8	10.5
73574_at			AI290214	4.8	3.7	8.1
62165_at	x 006 protein		W72231	4.8	3.2	5.9
40824_at	RAN binding protein 16	RANBP16	AB018288	4.8	2.5	5.3
50244_at	nucleotide binding protein 2 (E.coli MinD like)	NUBP2	AA552351	4.8	5.6	6.4
39971_at	lymphoblastic leukemia derived sequence 1	LYL1	M22637	4.8	3.5	4.8
32820_at	CCR4-NOT transcription complex, subunit 4	CNOT4	U71267	4.7	4	10.4
67963_at			AA436007	4.7	4.7	3.1
87874_at			T69727	4.7	4.4	8.2
44202_at			AA417797	4.7	3.8	6.1
52204_at			AA894564	4.7	5	4.3
74680_r_at	FLN29 gene product		AW002527	4.7	7	6.3

Figure 17AA

53670_at				AA664156	4.7	17.4	9.1
52855_at				AI885781	4.7	11.3	14.6
45254_at	translocase of inner mitochondrial membrane 10 (yeast) homolog		TIMM10	W02499	4.7	4.6	4.9
44705_at	hypothetical protein			AA133356	4.7	3.9	4.9
42831_at				AI201843	4.7	4.2	4.7
44823_s_at				AA521476	4.7	4.4	4.8
35166_at	Down syndrome critical region gene 3		DSCR3	D87343	4.6	7.1	5.7
37493_at				H04668	4.6	4.7	3.3
36953_at	MAD (mothers against decapentaplegic, Drosophila) homolog 4		MADH4	U44378	4.6	3.1	3.4
82032_at				AI823999	4.6	5.2	3.5
55362_at				AI653767	4.6	18.2	19.8
41749_at	ES1 (zebrafish) protein, human homolog of		C21ORF33	U53003	4.6	3	4.1
32562_at	endoglin (Osler-Rendu-Weber syndrome 1)		ENG	X72012	4.6	4.7	6
48810_at	hypothetical protein DKFZp761H221			AI547243	4.6	3.7	6.5
45758_at				AA034095	4.6	3	5.9
65620_at	hypothetical protein FLJ14005			W52855	4.6	4.4	4.7
45661_at				N90348	4.5	5.5	6.7
46234_at	CGI-148 protein			AA043242	4.5	4	4.9
32317_s_at	sulfotransferase family, cytosolic, 1A, phenol-preferring, member 2		SULT1A2	U34804	4.5	3.1	4.1
39135_at	KIAA0767 protein			AB018310	4.5	3.7	4.3
52081_at	ZAP3 protein			AI525457	4.5	3.3	4.3
32635_at	transcriptional intermediary factor 1 gamma			AB029036	4.4	3.7	6.1
56940_g_at				AI963304	4.4	3.6	4.3
49312_at	CGI-102 protein			AA747303	4.4	3.2	7.4
62495_f_at				AI433785	4.4	17.4	28.1
43489_at	hairy/enhancer-of-split related with YRPW motif-like		HEYL	AL040197	4.4	3.2	5.1
48992_at				AA435933	4.4	4.2	6.4
77840_f_at				AI720888	4.4	8.1	16.6
41865_at				AF052185	4.4	3.3	4.3
33916_at	imidazoline receptor candidate			AB023192	4.4	3.8	3.6
48094_at				AA531025	4.4	3.8	4.1
44085_at				AL040188	4.4	4	4.6

Figure 17BB

57261_f_at	ribosomal protein, large, P0	RPLP0	AA588862	4.4	4.1	7.9
39412_at	zinc finger protein 173	ZNF173	U09825	4.3	5	16.9
1448_at	proteasome (prosome, macropain) subunit, alpha type, 3	PSMA3	D00762	4.3	4	5.6
38899_s_at	hypothetical protein FLJ20693		U95822	4.3	5.7	3.1
40332_at	7-60 protein		AF109134	4.3	4.2	5.3
36899_at	special AT-rich sequence binding protein 1 (binds to nuclear matrix/scaffold-associating DNA's)	SATB1	M97287	4.3	4.8	6.6
56927_at			W49628	4.3	4.5	7.8
55361_at			AI821565	4.3	6.5	8.3
40848_g_at	KIAA0750 gene product		AB018293	4.3	4.2	4
35199_at	KIAA0982 protein		AB023199	4.3	2.7	4.4
33920_at	diaphanous (Drosophila, homolog) 1	DIAPH1	AF051782	4.3	3.2	3.8
38424_at	KIAA0747 protein		AB018290	4.3	2.9	4
210_at	phospholipase C, beta 2	PLCB2	M95678	4.3	3.7	3.6
53906_at	differentially expressed in FDCP (mouse homolog) 6	DEF6	AA523303	4.3	4.3	4.3
54158_at			AI453531	4.3	2.8	4.6
45115_at			AA909042	4.3	4.2	4.5
560_s_at			M63589	4.3	4.3	4.3
61067_f_at			AA470798	4.2	3.7	6.4
48315_at			AA286940	4.2	4.9	5.5
60690_at			AA203328	4.2	4.4	7.2
78732_f_at			AI094933	4.2	9.3	10.2
53848_at			AI422099	4.2	3.7	5.4
51713_at			AI660536	4.2	3.8	4.2
45285_at	proteasome (prosome, macropain) subunit, alpha type, 7	PSMA7	AA877820	4.2	3.4	5.1
81574_at	caspase recruitment domain protein 9		AI307612	4.2	3.6	4.2
52908_at	hypothetical protein FLJ21324		AA865619	4.2	3.9	4.1
62552_at			AA287801	4.2	3	3.3
44884_at			AA039908	4.1	7.3	6.8
2066_at	BCL2-associated X protein	BAX	L22474	4.1	4.6	9.8
41147_at			AF038186	4.1	15.4	3.6
37980_at	CBF1 interacting corepressor		U03644	4.1	9.1	8
55253_at			AA621047	4.1	3.8	4.3
82421_i_at			AI554417	4.1	6.2	8.8
59054_at			AA489100	4.1	9.9	4.8

Figure 17CC

80391_at	kinesin-like 2		KNSL2	AI953141	4.1	3.8	6.4
57660_r_at				R43504	4.1	3.4	5.6
38956_at	N-acetylglucosamine-1-phosphodiester alpha-N-acetylglucosaminidase			AF052111	4.1	3.1	3.9
38671_at	KIAA0620 protein			AB014520	4.1	4.3	5.3
35331_at	catenin (cadherin-associated protein), alpha-like 1		CTNNA1	U97067	4.1	3.3	4.1
48833_at	HSPC040 protein			AI749098	4.1	3.2	4.1
44078_at				W95017	4.1	5.1	5.5
48106_at	hypothetical protein FLJ20489			H14241	4.1	3.6	3.9
45335_at	CGI-28 protein			AI037884	4.1	2.7	4.4
52841_at				AA731740	4.1	4	4
46247_at	succinate-CoA ligase, GDP-forming, alpha subunit		SUCLG1	AA826261	4.1	3.3	5.5
54948_at				AI684747	4.1	3.5	3.9
77231_at				AI761622	4	3.1	7
63074_at				AA570454	4	3.7	4.7
57739_at	hypothetical protein FLJ20195			AI949010	4	3.5	8.7
65887_at	transducer of ERBB2, 2		TOB2	AI830948	4	3.4	5
35723_at	nudix (nucleoside diphosphate linked moiety X)-type motif 1		NUDT1	D16581	4	3.1	3
33569_at	macrophage lectin 2 (calcium dependent)			D50532	4	4	4
90328_at				AA771960	4	3.6	4
50926_s_at	fatty acid hydroxylase			R54585	4	4	4
64410_at	KIAA1150 protein			AA813014	4	2.6	3.6
54935_at	CD74 antigen (invariant polypeptide of major histocompatibility complex, class II antigen-associated)		CD74	N22918	4	3.2	3.7
32294_g_at	luteinizing hormone/choriogonadotropin receptor		LHCGR	M63108	3.9	3.2	7.5
34759_at				U68494	3.9	7.2	23.3
39150_at	ring finger protein 11		RNF11	U69559	3.9	3.2	6.1
51174_at	spinocerebellar ataxia 7 (olivopontocerebellar atrophy with retinal degeneration)		SCA7	AL043992	3.9	4.4	3.2
61191_at				AI539443	3.9	14.2	9.4
52770_at	LIS1-interacting protein NUDE1, rat homolog			AI857685	3.9	4.3	4.2
40266_at	KIAA1036 protein			AB028959	3.9	3.7	4.3
41504_s_at	v-maf musculoaponeurotic fibrosarcoma (avian) oncogene homolog		MAF	AF055376	3.9	3.2	4.1

Figure 17DD

52238_s_at	transforming, acidic coiled-coil containing protein 3	TACC3	AI990642	3.9	3.4	3.5
65813_at	zinc finger protein homologous to Zfp-36 in mouse	ZFP36	N32721	3.9	2.8	4.8
60863_s_at	hypothetical protein from EUROMAGE 2021883		AA526910	3.9	2.8	4.3
44343_s_at			AI074877	3.9	3.1	3.8
57210_at			AI862775	3.9	3.3	4
55564_at	spleen tyrosine kinase	SYK	AA521157	3.9	3.8	3.9
57222_at			AA203532	3.9	2.6	3.5
43991_at	hypothetical protein FLJ13576		W19970	3.9	2.8	3.9
47640_at	Nedd4-like ubiquitin-protein ligase		AI668780	3.9	4.1	5.1
38704_at	actin binding protein; macrophin (microfilament and actin filament cross-linker protein)		AB007934	3.8	3.6	4.7
68675_at	ficolin (collagen/fibrinogen domain-containing) 3 (Hakata antigen)	FCN3	AI652910	3.8	4.7	6.3
51053_at			AI122787	3.8	4.8	6.7
58827_at			AI992368	3.8	4.2	5.3
48767_at	ALEX1 protein		AI693923	3.8	5.8	16.6
67162_r_at			AI366683	3.8	3.3	3.8
38487_at	KIAA0246 protein		D87433	3.8	3.6	3.9
57223_at	potassium voltage-gated channel, Isk-related family, member 3	KCNE3	AI982768	3.8	3.2	3.5
44569_at			AA961420	3.8	3.6	3.6
54765_at	xeroderma pigmentosum, complementation group C	XPC	AL042673	3.8	2.8	4.2
46569_at			AA552976	3.8	2.8	3.8
40700_at	nuclear body protein Sp140		U36500	3.7	13.3	19
37468_at	Janus kinase 2 (a protein tyrosine kinase)	JAK2	AF058925	3.7	4	3.1
31671_at	RNA binding motif, single stranded interacting protein 1, pseudogene	RBMS1P	D82351	3.7	4.6	3.9
32628_at	zinc finger protein 161	ZNF161	D28118	3.7	3.5	7.2
35847_at	ubiquitin specific protease 24	USP24	AB028980	3.7	3	4.7
40390_at	serine dehydratase	SDS	J05037	3.7	5.2	16.9
38082_at	KIAA0650 protein		AB014550	3.7	3	12.1
58005_at	hypothetical protein		AI480253	3.7	6.7	3.8
74867_at			AI038997	3.7	4.3	3
65839_at	PC3-96 protein		AW022836	3.7	4.4	6.3
42804_at			W86160	3.7	5.1	5.4

Figure 17EE

81806_at	KIAA1681 protein								
54020_at	zinc finger protein 161	ZNF161							
55959_at									
46042_at									
48084_at									
80889_at									
47390_at									
65923_at									
61896_at									
43105_at									
44936_at									
72742_at									
36571_at	topoisomerase (DNA) II beta (180kD)								
90257_at	hypothetical protein DKFZp547D065	TOP2B							
51056_at	hypothetical protein DKFZp434D0412								
43992_at									
51163_at	hypothetical protein FLJ22405								
65465_at									
55678_at	hypothetical protein MGC10986								
80251_r_at	ATPase, Ca++ transporting, ubiquitous	ATP2A3							
87754_g_at	serologically defined colon cancer antigen 3	SDCCAG3							
38659_at	suppressor of clear, C. elegans, homolog of								
37558_at	IGF-II mRNA-binding protein 3								
81276_at									
43281_at									
49600_at									
90038_at	sin3-associated polypeptide, 18kD	SAP18							
87221_at									
58622_at									
70323_f_at									
1427_g_at	Src-like-adaptor	SLA							
56238_at	PDZ-LIM protein mystique								
57461_at									
32259_at	enhancer of zeste (Drosophila) homolog 1	EZH1							
43850_s_at	PP1201 protein								

Figure 17FF

58564_s_at				N78066	3.4	3.2	3.5
38459_g_at	cytochrome b-5		CYB5	L39945	3.4	6.8	5.6
40244_s_at	hypothetical protein FLJ11585			AI743654	3.4	2.6	3.5
40745_at	adaptor-related protein complex 1, beta 1 subunit		AP1B1	L13939	3.4	2.7	3.2
58498_at	KIAA1378 protein			AI082236	3.4	2.5	3.4
80436_at				AI935644	3.4	3.3	4
45276_g_at	protein x 0001			W30943	3.4	3	4.9
57156_at	HSPC023 protein			W92964	3.4	3.2	4.9
45543_r_at	mitochondrial import receptor Tom22			AI359916	3.4	3.4	2.8
54483_at				AA760777	3.4	3.4	3.4
34678_at	fer-1 (C.elegans)-like 3 (myoferlin)		FER1L3	AL096713	3.3	8.2	3.4
32455_s_at	proline and glutamic acid rich nuclear protein			U88153	3.3	2.6	3.2
479_at	disabled (Drosophila) homolog 2 (mitogen-responsive phosphoprotein)		DAB2	U53446	3.3	3	3.4
90610_at	leucine-rich neuronal protein		LRN	AI654857	3.3	2.5	4.3
49463_at	suppressor of fused			AI760042	3.3	3.1	3
44047_at	hypothetical protein FLJ20160			AA133311	3.3	3.3	3.5
52932_at	kangai 1 (suppression of tumorigenicity 6, prostate; CD82 antigen (R2 leukocyte antigen, antigen detected by monoclonal and antibody IA4))		KAI1	W51778	3.3	2.9	2.6
45339_at	CGI-39 protein; cell death-regulatory protein GRIM19			AA181145	3.3	3.7	4.4
53940_at				AI953658	3.3	3.1	3.4
46220_at	peroxisomal short-chain alcohol dehydrogenase			AI131178	3.3	3	3.7
32859_at	signal transducer and activator of transcription 1, 91kD		STAT1	M97935	3.2	18	8.2
38091_at	lectin, galactoside-binding, soluble, 9 (galectin 9)		LGALS9	Z49107	3.2	7.7	6.7
57737_at				AI150441	3.2	5.1	3.4
84320_at				AA602573	3.2	3.5	4.8
50081_at				AI765698	3.2	4.6	3.4
39774_at	oxidase (cytochrome c) assembly 1-like		OXA1L	X80695	3.2	3.1	5.4
39672_at	protein tyrosine phosphatase, non-receptor type 7		PTPN7	M64322	3.2	4.4	4.6
40296_at	X-prolyl aminopeptidase (aminopeptidase P) 2, membrane-bound		XPNPEP2	AL023653	3.2	2.7	3.2
47463_at	eukaryotic translation initiation factor 2C, 1		EIF2C1	AA453723	3.2	3.1	2.9
54865_at	coronin, actin-binding protein, 1C		CORO1C	N27805	3.2	2.9	3.2
60471_at	hypothetical protein FLJ22439			AA625133	3.2	2.9	3.1

Figure 17GG

43926_at				N36085	3.2	2.7	3.2
53956_at				AA449296	3.2	2.6	2.8
61119_at	CCAAT/enhancer binding protein (C/EBP), alpha	CEBPA		AI971171	3.2	2.7	3.1
38688_at	KIAA0461 protein			AB007930	3.1	3.3	7.4
68194_at				AA291194	3.1	4.8	10
88143_at				AI253134	3.1	3	4.1
64103_at				N22751	3.1	7.2	9.1
62480_at				AI446414	3.1	3.8	12.7
1933_g_at	ATP-binding cassette, sub-family C (CFTR/MRP), member 5	ABCC5		U83661	3.1	2.5	3
38976_at	coronin, actin-binding protein, 1A	CORO1A		D44497	3.1	3.1	3
41600_at	proliferation-associated 2G4, 38kD	PA2G4		U59435	3.1	2.9	3
57030_at				W26295	3.1	3.1	3.1
62889_at	KIAA0675 gene product			D29070	3.1	3.1	3.1
63600_f_at				AW021173	3.1	2.9	3.2
65299_at				C15819	3.1	3.3	3.1
36118_at	nuclear receptor coactivator 1	NCOA1		AJ000882	3	4.5	4.4
54487_at				AA044626	3	3.1	5.1
40024_at	src homology three (SH3) and cysteine rich domain	STAC		D86640	3	3.1	3.3
36792_at	tropomyosin 1 (alpha)	TPM1		Z24727	3	3	2.9
38336_at	KIAA1013 protein			AB023230	3	2.5	2.9
48595_at	hypothetical protein FLJ10210			AA707400	3	2.8	2.8
32800_at	retinoid X receptor, alpha			U66306	3	2.9	2.6
65612_at	hypothetical protein MGC3180			AW006145	3	3	3
43854_at				AW004730	3	2.8	2.9
43499_at	tropomyosin 1 (alpha)	TPM1		N76371	3	2.6	2.6
35342_at				AF052159	2.9	2.5	2.6
36135_at	nucleolar protein p40; homolog of yeast EBNA1-binding protein			U86602	2.9	2.7	2.9
40695_at	IMP (inosine monophosphate) dehydrogenase 1	IMPDH1		J05272	2.9	2.7	3.7
38770_at	a disintegrin and metalloproteinase domain 12 (meltrin alpha)	ADAM12		AF023476	2.9	2.9	2.7
38290_at	regulator of G-protein signalling 14	RGS14		AF037195	2.9	2.6	2.7
78442_at	hypothetical protein MGC4692			AI819340	2.9	2.9	2.9
49343_at				AI819630	2.9	2.9	3.2

Figure 17HH

44849_at	beta-site APP-cleaving enzyme	BACE	AI356651	2.9	2.9	2.9
55671_at	FH1/FH2 domain-containing protein		N90883	2.9	2.7	2.8
35338_at	paired basic amino acid cleaving enzyme (furin, membrane associated receptor protein)	PACE	X17094	2.8	3.1	3.5
59452_at			AI719167	2.8	2.8	2.9
57918_at	DKFZP586G1517 protein		AA779617	2.8	2.6	2.8
60493_at			AI344328	2.7	2.5	2.6
59504_at			AA932206	2.6	2.6	2.5
49349_at	PP2135 protein		W72919	2.6	2.6	2.7
85006_at			AA601045	2.6	2.6	2.6
44143_at	McKusick-Kaufman syndrome	MKKS	AA399076	2.6	3.1	3.5
57346_at	Mad4 homolog		AA831438	2.5	3	3.6
88767_at			AW008519	2.5	2.5	2.5
91825_at			AI375222	2.5	2.5	2.5
50198_at			AI208272	2.5	2.5	2.5
47627_at	cytochrome P540, subfamily IIS, polypeptide 1		AI445492	2.5	2.5	2.5

Figure 17II